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Result
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence:
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Perfect score:
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Maximum
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   92:
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1691
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first 45 summaries
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Abj04079
Aae21296
Adi16993
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81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.2	81.3	81.3	81.3	81.7	82.5	82.5
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Adh08520 Mrs	Ad196466 Hur	Abw00803 Hur			Abp96696 Hui		Adil6636 Hu	Adil6991 Hu	Aae21288 Hu	Abp95617 Hu	Aae17074 Hu	Aa:104371 Hui		Aag64294 Hui	Adc17728 Hui	Aay87664 Hu		Aay30162 Hu	Adw02566 Hur	Aay30161 Hu
MrgX1.	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human	Human
ω	e G	GPC	GPC	Dor	g G	Nov	NOV	NOV.	PIM	GPC	G-p	G-b	G ဗ	GTP	TGR	a ဗ	GPC	dor	sen	dor

ALIGNMENTS

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AC AAX3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
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(ASTR) ASTRA ASTRA PHARMA INC. AB.

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WPI; 199 N-PSDB; 1999-405162/34. DB; AAZ10067.

and human dorsal root receptors and related polynucleotides, useful identifying agents for anesthesia and analgesia.

Claim 6; Page 39-41; 72pp; English.

This is the human dorsal root receptor 1 (hDRR1) protein sequence. It is a G protein coupled receptor that is expressed preferentially in dorsal root ganglia. hDRR1 can be used to create antibodies against hDRR1. The dorsal root ganglia area of the central nervous system (c is densely innervated with primary or afferent neurons involved in transmission, modulation and sensation of pain. The DRR's which are expressed in this region of the CNS may be used for assays for the identification of new agents for anaesthesia and analgesia. (Updated 20-MAR-2003 to correct PA field.) (Updated (CNS) This 9

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sensory neuron-specific G protein-coupled receptor 1; SNSR1;
g-protein coupled receptor; gpcr; diagnosis; cardiovascular disease;
gastrointestinal disease; liver disease; cancer; neoplasm; inflammation;
hematological disease; respiratory disease; neurological disease;
genitourinary disease; cardiovascular-gen.; cardiant; vasotropic;
antiarrhythmic; antiarceriosclerotic; hypotensive; gastrointestinal-gen.;
antiinflammatory; antiulcer; hepatotropic; antiallergic; dermatological;
thyrominetic; immunosuppressive; antianemic; cytostatic; hemostatic;
antiasthmatic; respiratory-gen.; cns-gen.; antiparkinsonian; nootropic;
neuroprotective; cerebroprotective; nephrotropic; uropathic; receptor.
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                           Screening for therapeutic agents, useful for treating e.g., cardiovascular disorders, comprises contacting a test compound with sensory neuron-specific G protein-coupled receptor 1 (SNSR1) polyper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADW46672 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                      04-JUN-2004; 2004WO-EP006078
   detecting binding
                                                                                                                                                  2005-057894/06.
DB; ADW46671.
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Pred. No. 2.4e-175;
; Mismatches 0;
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Disclosure; SEQ ID NO 2; 122pp; English

The invention relates to a method of screening for therapeutic agents C treating diseases associated with sensory neuron-specific G protein-C coupled receptor (SNSR1). The method comprises contacting a test compound with SNSR1 polypeptide or polynucleotide and detecting the compound with SNSR1 polypeptide or polynucleotide and detecting the compound with SNSR1 polypeptide or polynucleotide, or different concentration, (b) in the absence of the test compound, or (c) at a contentration of the test compound. Also described are (i) a method of diagnosing a disease defined above in a mammal, (ii) a pharmaceutical composition for the treatment of the disease above a comprising the SNSR1 polypeptide, (iii) a SNSR1 polypeptide or polynucleotide, or a cherpeutic agent which binds to the SNSR1 polypeptide or which regulates the SNSR1 polypeptide activity such as a small molecule, an RNA molecule, an antisense oligonucleotide, a polypeptide, an antibody, or a ribozyme, and (iv) a method for the preparation of a pharmaceutical composition or useful for preparing a pharmaceutical composition or treating the above diseases. The SNSR1 regulatory compounds are useful for preparing a pharmaceutical composition for treating condiscores, respiratory diseases, neurological disorders, respiratory diseases, hematological disorders, respiratory diseases, mentological disorders, respiratory diseases include antipological disorders, and chronic pancreatitis. Inflammatory diseases, or unological disorders, and chronic pancreatitis. Inflammatory diseases include atherosclerosis tiver diseases include jaundice, Crijter-Najjar, chashing the disease. Cardiovascular diseases include atherosclerosis, stroke, and Alabelmer's metaplasia, stress gastritis, gastric diseases, and chronic pancreatitis. Inflammatory diseases include atopic diseases, and chronic pancreatitis. Inflammatory diseases include atopic diseases, and chronic pancreatitis. Inflammatory diseases include atopic diseases, and chronic pancreatitis. Inflammatory diseases include a

Sequence 322 AA;

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                                                                                                                                                                                                                                                          1 MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                    VSIYILNLVAANFLFLSGHIIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
                                                                                                                                               CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
                                                                                                                                                                                                   VSIYILNLVAANFLFLSGHIIFSPLFLINIRHPISKILSPVMTFPYFIGLSMLSAISTER 120
                                                                                                                                                                                                                                        MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
                                      RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                           CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
VDEGGGWLPQETLELSGSKLEQ
                         RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                              100.0%; Score 1691; DB 9; 100.0%; Pred. No. 2.4e-175; tive 0; Mismatches 0;
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VDEGGGWLPQETLELSGSKLEQ

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RESULT 3
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Best Local (
                                                                                                                                                                                                                     Matches 314;
                                                                                                                                                                                                                                                                                                     The present sequence represents a human-derived G protein-coupled protein designated hHI77213V, which is isolated from the human hippocampus. The G protein-coupled receptor can be used for preventing, treating and diagnosing genetic diseases associated with G protein-coupled protein, and for regulating cellular functions. The protein can be used to prevent and treat disorders associated with G protein-coupled protein dysfunction. It can also be used to identify G protein-coupled protein ligands and generating antibodies and antisers against the protein. It is also useful in constructing recombinant receptor protein expression systems, developing receptor-binding assay systems and screening drug candidates, and can be used as a probe in the genetic diagnosis of G protein-coupled protein disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human-derived G protein-coupled e.g. in determining ligands and dysfunction of the protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; G protein-coupled receptor; hippocampus; diagnosis; screening; genetic disease; cellular function regulation.
                                                                                                                                                                                                                                                                               Sequence 322 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2; Page 92-93; 97pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB;
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                                                                                                                                                                                                                     Local Similarity 97.
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                                                          CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
                                                                                                                    VSIYILNLVAANFLFLSGHIIFSPLFLINIRHPISKILSPVMTFPYFIGLSMLSAISTER 120
                                                                                                                                                                          MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA 60
                IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
                                        CLSILMPIWYHCRRPRYLSSVMCVLLWALSLLRGILEMMFCDFLFSGANSVWCETSDFIT
                                                                                                                                                           MDSTIPVLGTELTPINGREETPCYKQTLSFTGLTCIVSLVALTGNAVVLWLLGCRMRRNA
                                                                                                    VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
Terao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                  Score 1647;
Pred. No. 1.
                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein and encoding nucleic acid, useful treatment of diseases associated with
                                                                                                                                                                                                                                  DB 3;
L.5e-170;
                                                                                                                                                                                                                                               Length 322;
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RESULT 4
AD116994
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27-FSB-2001;
27-FSB-2001;
02-MAR-2001;
02-MAR-2001;
14-MAR-2001;
14-MAR-2001;
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08-FEB-2001;
09-FEB-2001;
15-FEB-2001;
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31-JAN-2001;
31-JAN-2001;
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16-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; NOVX; cardiomyopathy; atherosclerosis; cancer; diabetes; inflammation; autoimmune disorder; allergy; blood disorder; acquired immunodeficiency syndrome; AIDS; obesity; asthma; immunoglobulin (Ig)A nephropathy; cirrhosis; arthritis; Alzheimer's disease; infection; str.
                                                                                                                                                                                                                                                                                                                                                                                                                                 14-MAR-2001
14-MAR-2001
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15-MAR-2001;
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2001US-0265514P.

2001US-026551P.

2001US-0266406P.

2001US-0266767P.

2001US-0266975P.

2001US-0266975P.

2001US-0267057P.

2001US-0267057P.
2001US-0276448P.
2001US-0276450P.
2001US-027639P.
2001US-0276768P.
2001US-0278775P.
2001US-0278775P.
2001US-0278775P.
2001US-027884P.
2001US-028014P.
2001US-028014P.
2001US-0285133P.
2001US-0285133P.
2001US-0285133P.
2001US-0285137P.
2001US-028532P.
2001US-0288532P.
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2001US-0288532P.
2001US-0288532P.
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2001US-02687974P
2001US-0271664P
2001US-0271839P
2001US-0271855P
2001US-02727889
2001US-0272784P
2001US-0275947P
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2001US-0275949P
2001US-0275949P
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                                                                                                                                                                                                                                                                                           Cd disgnostic and prognostic assays and furthermore in the treatment of cdiverse pathological conditions. The present invention describes novel conditions as well as methods to modulate their cc expression using antisense oligos, ribozymes and peptide muclaic acids. The NOVX polypeptides, polynuclectides and antibodies are useful in cc treating or preventing NOVX-associated disorders, e.g. cardiomyopathy, cc atherosclerosis, cancer and diabetes. Furthermore, they may be used in cc treating or preventing diseases such as inflammation, autoimmune cd disorders, allergies, blood disorders, acquired immunodeficiency syndrome (AIDS), obesity, asthma, immunoglobulin (Ig)A nephropathy, cirrhosis, carthritis, Alzheimer's disease, infections, stroke, muscular dystrophy and epilepsy. Accordingly, these molecules have many activities including cytostatic, cardiant. antinflammatory, immunosuppressive, antiallergic, antiastatic, nephrotropic, antiastratic, hepatotropic, cantiastratic, hepatotropic, conversative, nootropic, antiastratic, hepatotropic, antiallergic, antiastratic, hepatotropic, antiastratic, nephrotropic, antiastratic, hepatotropic, antiastratic, for example, conserved to dentify small molecules that modulate or inhibit, for example, conserved to dentify small molecules that modulate or inhibit, for example, conserved to dentify small molecules that modulate or inhibit, for example, conserved to dentify small molecules that modulate or inhibit, for example, and incording and anyticalles that modulate or inhibit, for example, conserved to the invention.
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28-AUG-2001;
31-AUG-2001;
07-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to a novel nucleic acids, and encoded polypeptides thereof, which have properties related to the stimulation of biochemical or physiological responses in a cell, tissue, organ or organism. Specifically, it refers to the use of biologically active fragments for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-SEP-2001;
12-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating NOVX-associ atherosclerosis, or pharmacogenomics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tchernev VT, Spytek KA, Zerhusen BD, Patturajan M, Shimkets RA;
Li L, Gangolli EA, Padigaru M, McCeson DW, Rastelli L, Miller CE;
Gerlach VL, Taupier RJ, Gusev VY, Colman SD, Wolenc AR, Pena CEA;
Furtak K, Grosse WM, Alsobrook JP, Lepley DM, Rieger DK, Burgess CE;
                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 530; 1498pp;
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18-OCT-2001;
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13
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                                                                          VSIYILNLVAANFLFLSGHIIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
  VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
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2001US-031202DP.
2001US-0312889P.
2001US-031390P.
2001US-0315470P.
2001US-0316447P.
2001US-031811SP.
2001US-031811SP.
2001US-03181290P.
2001US-03181290P.
2001US-03181290P.
2001US-0318129P.
2001US-032379P.
2001US-032379P.
2001US-0333308P.
2001US-0333701P.
                                                                                                                                                            Conservative
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                                                                                                                                                                              97.4%;
                                                                                                                                                            4.
                                                                                                                                                            Score 1647; DB 5;
Pred. No. 1.5e-170;
4; Mismatches 4;
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RESULT 5
ADI1 6999
ID 10 15-3
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XX ADI1
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XX ADI1
XX ADI1
XX Infil
XX huma
XX huma
XX huma
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07-FEB-2001

08-FEB-2001

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14-MAR-2001

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07-FEB-2001;
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  2001US-0266767P
2001US-0266767P
2001US-026795P
2001US-0268974P
2001US-02786974P
2001US-0271664P
2001US-0271859P
2001US-0271859P
2001US-0273046P
2001US-0275947P
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2001US-0275947P
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2001US-0276448P
2001US-0276450P
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2001US-0276479P
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2001US-0276479P
2001US-0278779P
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2001US-0279882P
2001US-0279882P
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2001US-0266406P.
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13-AUG-2001

16-AUG-2001

16-AUG-2001

16-AUG-2001

21-AUG-2001

28-AUG-2001

31-AUG-2001

07-SEP-2001

07-SEP-2001

12-SEP-2001
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19-JUN-2001;
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2001US-029324P.
2001US-0312020P.
2001US-0312889P.
2001US-0313390P.
2001US-0315470P.
2001US-0318118P.
2001US-0318118P.
2001US-0318118P.
2001US-0318118P.
2001US-0318149P.
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2001US-0318149P.
2001US-0318149P.
2001US-031839P.
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2001US-0285133P.
2001US-0285749P.
2001US-0288327P.
2001US-0288327P.
2001US-029404P.
2001US-0294473P.
2001US-0294473P.
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Gerlach VL, Taup--Gerlach VL, Grosse WM, ey VT, Spytek KA, Zerm.
Gangolli EA, Padigaru M, Anderen.
Gangolli EA, Padigaru M, Colman SD,
NU, Taupier RJ, Gusev VY, Colman SD,
NU, Taupier RJ, Gusev VY, Lepley DM, Patturajan M, Shimkets RA; son DW, Rastelli L, Miller CE; lman SD, Wolenc AR, Pena CEA; Rieger DK, Burgess CE;

New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing pharmacogenomics. 유

Disclosure; SEQ ID NO 535; 1498pp; English.

CC diagnostic and prognostic assays and furthermore in the treatment of CC diverse pathological conditions. The present invention describes novel human and murine NOVX protectins, as well as methods to modulate their CC expression using antisense oligos, ribozymes and peptide nucleic acids. The NOVX polypeptides, polynucleotides and antibodies are useful in CC treating or preventing NOVX-associated disorders, e.g. cardiomyopathy, CC atherosclerosis, cancer and diabetes. Furthermore, they may be used in CC disorders, allergies, blood disorders, acquired immunodeficiency syndrome CC disorders, allergies, blood disorders, acquired immunodeficiency syndrome CC disorders, Alzheimer's disease, infections, stroke, muscular dystrophy CC arthritis, Alzheimer's disease, infections, stroke, muscular dystrophy CC arthritis, Alzheimer's disease, infections, stroke, muscular dystrophy CC arthritis, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, antiasteriat, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, CC haemostatic, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, continuous contropic, antibacterial, virucide, antiparasitic, neurogrotective, nootropic, antibacterial, virucide, antiparasitic, consucration, continuous contropic, antibacterial, virucide, antiparasitic, consucration, continuous contropic, antibacterial, virucide, antiparasitic, consucration, consucration, continuous contropic, antiparasitic, hepatotropic, antiparasitic, hepatotropic, antiparasitic, assays to identify small molecules that modulate or inhibit, for example, continuous contropic, antiparasitic, antiparasit This invention relates to a novel nucleic acids, and encoded polypeptides thereof, which have properties related to the stimulation of biochemical or physiological responses in a cell, tissue, organ or organism. Specifically, it refers to the use of biologically active fragments for

Sequence 322 8

Best Local Sin Matches 314; Local Similarity Conservative 97.4**%**; 97.5**%**; 4 Score 1647; Pred. No. 1 Mismatches .5e-170; DB 5; Length 322; ٥, Gaps

> ADCORPTION (NAAD-) (ADSC-) 18-JUN-2001; 2001JP-00246789 18-JUN-2002; 2002EP-00013517 02-JAN-2003 EP1270724-A2 human; GPCR; guanosine triphosphate-binding protein coupled receptor; Human GPCR protein SEQ ID NO:1274. 01-JAN-2004 ADC86821; ADC86821 standard; σ 2003-315783/31. DB; ADC86820. therapy 301 241 241 181 181 121 121 61 61 <u>ــر</u> NAT INST ADVANCED IND SCI & TECHNOLOGY.
> CENT ADVANCED SCI & TECHNOLOGY INCUBATI VDEGGGWLPQETLELSGSKLEQ RIHLDWKYLFCHVHLYSIFLSALNSSANPIIYFFYGSFRQRQNRQNLKLYLQRALQDTPE IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER VSIYILNLVAANFLFLSGHIIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE VDEGGGWLPQETLELSGSRLEQ IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS (first Akiyama protein; entry) Ķ 322 A 322 322 INCUBATIO 300 240 180 180 120 120 60

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片

New polynucleotide, useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor.

Claim 2 SEQ ID NO 1274; 28pp; English.

The invention relates to a novel polynucleotide encoding a guanosine triphosphate-binding protein coupled receptor (GPCR). A polynucleotide of the invention may have a use in gene therapy. The polynucleotide and polypeptide are useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor. The protein sequences shown in ADC85549-ADC87617 represent GPCR's of the invention.

Sequence 322 AA;

Query Match

0,

97.4%; Score 1647; DB 7; Length 322,

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RESULT 7
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XX AD04
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CCC AD04
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CCC AD16
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The invention relates to a non-human mammal that carries a DNA integrated with a foreign HI7T213 or its mutant gene, or a part of it. The non-human animal is particularly a rat. Such gene shows phenotypes of e.g. cataract onset, transient skin rash and proliferation-promoting activity. The foreign HI7T213 gene is a gene that encodes a G protein-coupled receptor (GPCR) protein HI7T213. The protein, its encoded DNA and constructed transgenic animals are useful for clarifying pathological mechanism,
                                                                                                                                                                                                                                                                       pathological mechanism, preventives or remedies
                                                                                                                                                                                                  Claim 3; SEQ ID NO 3; 161pp; Japanese.
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ic; nephrotropic; antiinflammatory; dermatological; analgesic;
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                                                                                                                                                                                                                                                                       encoded DNA and transgenic animals for clarifying hanism, developing therapeutic methods and screening emedies for related diseases e.g. cataract, cancer, a
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4; Mismatches 4;
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Claim

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Page 90-91; 97pp;

Japanese

The present sequence represents a human-derived designated hH17T213, which is isolated from the

G protein-coupled human hippocampus

Drotein

Human-derived G protein-coupled e.g. in determining ligands and dysfunction of the protein.

protein and encoding nucleic acid treatment of diseases associated

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Best Local Simi
Matches 314;
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                                                                                                                                                                                                                                    Human; G protein-coupled receptor; hippocampus; genetic disease; cellular function regulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 322
                                                                                                                                                                                                                                                              Human G protein-coupled receptor hHI7T213
                                                                                                                                                                                                                                                                                                                  AAY90761 standard;
                                                                                N-PSDB;
                                                                                                          Watanabe
                                                                                                                                             01-OCT-1998;
                                                                                                                                                              30-SEP-1999;
                                                                                                                                                                                 13-APR-2000
                                                                                                                                                                                                  WO200020455-A1
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Best Local S
Matches 313
                     27-NOV-2000;
12-DEC-2000;
20-FEB-2001;
20-FEB-2001;
06-APR-2001;
06-APR-2001;
06-APR-2001;
14-MAY-2001;
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                                                                                                                                                                                                                                                                                     26-NOV-2001; 2001WO-US044386
                                                                                                                                                                                                                                                                                                                                       30-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; G-protein coupled receptor; GPCR; hRUP28; hRUP29; hRUP30; hRUP31; hRUP32; hRUP33; hRUP35; hRUP36; hRUP37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human G protein coupled receptor hRUP37.
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5; Mismatches 4;
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Best Local S
Matches 313
                                                                                                                                 Human; mas-related gene; G-protein coupled receptor; receptor; sensory perception; pain; analgesic; MrgX3
 04-MAY-2000; 2000US-0202027P
                         04-MAY-2001; 2001WO-US014519.
                                                                                                                                                                        Human MrgX3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences of several human G-protein coupled receptors (GPCRs). These can be use the identification of candidate compounds as receptor agonists or agonists having applicability as therapeutic agents. The present is a GPCR protein of the invention
                                                                                                                                                                                                    01-JUL-2002
                                                                                                                                                                                                                               AAE21296;
                                                                                                                                                                                                                                                        AAE21296 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel endogenous and non-endogenous versions of G protein-coupled receptor useful for identification of candidate compounds as receptorals or antagonists for use as therapeutic agents.
                                                                               WO200183555-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 322 AA;
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97.2%;
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Pred. No. 5.2e
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03-NOV-2000;
19-APR-2001;
                        human; NOVX; cardiomyopathy; atherosclerosis; cancer; diabetes; anflammation; autoimmune disorder; allergy; blood disorder; acquired immunodeficiency syndrome; AIDS; obesity; asthma; immunoglobulin (Ig)A nephropathy; cirrhosis; arthritis; Alzheimer's disease; infection; str.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to Mrg (mas-related gene) protein, which is a G-protein coupled receptor and drg-12 protein, which is a receptor. The invention is useful for identifying compounds that bind to it, especially agonists or antagonists. Administration of an agent (e.g. the identified agonist) that increases the expression of Mrg in a mammal may be used for treating impaired sensory perception in a mammal, especially pain. The antagonist may also be useful for treating impaired sensory perception in a mammal. The present sequence is human MrgX3 protein
                                                                                                                                                                      Human
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2001US-0285493P.
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5.2e-170;
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02-FEB-2001
07-FEB-2001
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08-FEB-2001
19-FEB-2001
19-FEB-2001
27-FEB-2001
27-FEB-2001
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11-FARR-2001
11-MAR-2001
11-MA
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                                                ev VT, Spytek KA, Zerhusen BI
Gangolli EA, Padigaru M, And
h VL, Taupier RJ, Gusev VY,
K, Grosse WM, Alsobrook JP,
                                                                                                                     CURAGEN CORP
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2001US-029324P
2001US-0312020P
2001US-0312889P
2001US-0313908P
2001US-0313490P
2001US-0315447P
2001US-0318118P
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2001US-0318149P
2001US-032379P
2001US-032379P
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                                                                                          Zerhusen BD,
                                                                            Anderson DW,
                                                                                          Patturajan M,
                                                    ıjan M, Shimkets RA;
Rastelli L, Miller CE;
Wolenc AR, Pena CEA;
1, Rieger DK, Burgess C
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NOVX polypeptides 2002-706998/76.

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nucleic

acids, useful

for preventing

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Colman SD, Lepley DM,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to a novel nucleic acids, and encoded polypeptides CC thereof, which have properties related to the stimulation of biochemical cor physiological responses in a cell, tissue, organ or organism. CC Specifically, it refers to the use of biologically active fragments for CC diverse pathological conditions. The present invention describes novel thuman and murine NOVX proteins, as well as methods to modulate their CC The NOVX polypeptides, polymorleotides and antibodies are useful in CC treating or preventing NOVX-associated disorders, e.g. cardiomyopathy, CC treating or preventing diseases such as inflammation, autoimnune CC disorders, allergies, blood disorders, acquired immunodeficiency syndrome (AIDS), obesity, asthma, immunoglobulin (Ig)A nephropathy, cirrhosis, cardiers, allergies, blood disorders, acquired immunodeficiency syndrome (AIDS), obesity, asthma, immunoglobulin (Ig)A nephropathy, cirrhosis, cardiant, antiinflammatory, immunosuppressive, antialergic, and policys, Accordingly, these molecules have many activities including cytostatic, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, neuroprotective, nootropic, antiarthritic, hepatotropic, antiallergic, consists and anticonvulsant. In addition, they are useful in screening convention, they are useful in screening conventions. This polypeptide is a homologue of a human NOVX protein cof the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Human NOVX protein homologue SeqID 534.
                                                                                                   ADI16998 standard;
                                    15-APR-2004
                                                                     ADI16998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pharmacogenomics.
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                                                                                                                                                                                                      VDEGGGWLPQETLELSGSKLEQ 322
                                                                                                                                                                                                                                                     RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                                                                                                                                                                                                                     RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
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                                                                                                                                                                                   VDEGGGWLPQETLELSGSRLEQ 322
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                                 (first entry)
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                                                                                                   protein; 322
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Pred. No. 5
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.2e-170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 322;
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114-MAR-2001;
15-MAR-2001;
15-MAR-2001;
16-MAR-2001;
26-MAR-2001;
26-MAR-2001;
26-MAR-2001;
29-MAR-2001;
29-MAR-2001;
29-MAR-2001;
11-APR-2001;
11-A
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08-FEB-2001

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27-FEB-2001
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31-JAN-2001;
31-JAN-2001;
31-JAN-2001;
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                                      (CURA-)
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                                      CURAGEN CORP.
                                                                    2001US-027984P

2001US-027984P

2001US-027984P

2001US-028993P

2001US-0283083P

2001US-0283083P

2001US-0285133P

2001US-0285749P

2001US-028832P

2001US-028832P

2001US-0298504P

2001US-0298473P

2001US-0299473P

2001US-039324P

2001US-0312020P

2001US-0312020P

2001US-0312932P

2001US-03129319

2001US-03129319

2001US-0313390P

2001US-031874P

2001US-0318115P

2001US-0318115P

2001US-0313379P

2001US-0333379P

2001US-033330308P

2001US-03333701P
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2001US-0271839P.
2001US-0272785P.
2001US-0275925P.
2001US-0275925P.
2001US-0275947P.
2001US-0275949P.
2001US-0276449P.
2001US-0276489P.
2001US-0276450P.
2001US-027676P.
2001US-0276852P.
2001US-0278652P.
2001US-0278652P.
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2001US-0267459P.
2001US-0267823P.
2001US-0268974P.
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2001US-0266767P.
2001US-0266975P.
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Spytek KA,

Zerhusen BD,

Patturajan M,

Shimkets RA;

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New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing
                                                                                                                        Furtak K,
                                                                                                                                              Gerlach
                                                                                                       n VI., Taur K., Grosse WM,
                                                                                                                    Taupier RJ, Gusev VY, Colman SD, rosse WM, Alsobrook JP, Lepley DM,
                                                                                                                                              Padigaru M, Ar
RJ, Gusev VY,
                                                                                                                                                                 Anderson DW,
                                                                                                                      Rastelli L,
Wolenc AR,
Molenc AR,
                                                                                                                                                Pena CEA;
                                                                                                                                                                   Miller CE;
                                                                                                                          Burgess CE;
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SEQ ID NO 534; 1498pp; English

pharmacogenomics.

CC or physiological responses in a cell, tissue, organism.

CC Specifically, it refers to the use of biologically active fragments for CC diverse pathological conditions. The present invention describes novel thuman and murine NOVX proteins, as well as methods to modulate their CC expression using antisense oligos, ribozymes and peptide nucleic acids. CC treating or preventing NOVX-associated disorders, e.g. cardiomyopathy, catherosclerosis, cancer and diabetes. Furthermore, they may be used in CC treating or preventing diseases such as inflammation, autoimmune CC disorders, allergies, blood disorders, acquired immunodeficiency syndrome CC (AIDS), obesity, asthma, immunoglobulin (Ig)A nephropathy, cirrhosis, cardiant, antiinflammatory, immunosuppressive, antiallergic, cardiant, antiinflammatory, immunosuppressive, antiallergic, cantiasthmatic, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, antiasthmatic, nephrotropic, antiarthritic, heparotropic, an invention relates to a novel nucleic acids, and encoded sof, which have properties related to the stimulation of invention polypeptides biochemical

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                                                                                         VSIYILMLVAANFLFLSGHIIFSDLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTFE
                            IAWLVELCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
                                                   CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGADSVWCETSDFIT
                                                                                VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
                                                                                                             MDSTIPVLGTELTPINGREETPCYKQTLSFTGLTCIVSLVALTGNAVVLWLLGCRMRRNA
                                                                                                                     MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
                                                                                                                                          Conservative
                                                                                                                                                 97.1%;
97.2%;
                                                                                                                                           <u>ن</u>
                                                                                                                                           Score 1642; DB 5;
Pred. No. 5.2e-170;
5; Mismatches 4;
                                                                                                                                                         Length 322;
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                                                                                                                                            Gaps
        300
                                                                   180
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                       240
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ABP81750 standard; protein; 322

04-MAR-2003 (first entry

Human G protein-coupled receptor MrgX3 protein SEQ ID NO:674

og R

g protein-coupled receptor; GPCR; antigenic peptide; gene therapy;
g protein-coupled receptor modulator; antibody; immune-related disease;
growth-related disease; cell regeneration-related disease; AIDS; cancer;
immunological-related cell proliferative disease; autoimmune disease;
Alzheimer's disease; atherosclerosis; infection; osteoarthritis; allergy;
osteoporosis; cardiomyopathy; inflammation, Crohn's disease; diabetes;
graft versus host disease; Parkinson's disease; multiple sclerosis; pain;
psoriasis; anxiety; depression; schizophrenia; dementia; memory loss;
mental retardation; epilepsy; asthma; tuberculosis; obesity; nausea;
hypertension; hypotension; renal disorder; rheumatoid arthritis; trauma; receptor; GPCR; antigenic peptide; gene therapy;

Homo sapiens.

WO200261087-A2

08-AUG-2002.

19-DEC-2001; 2001WO-US050107.

19-DEC-2000; 2000US-0257144P.

LIFESPAN BIOSCIENCES INC

GC, Roush CL, Brown JP

2003-046718/04

New isolated antigenic peptides e.g., for G protein-coupled receptors (GPCR), useful for diagnosing and designing drugs for treating conditions in which GPCRs are involved, e.g. AIDS, Alzheimer's disease, cancer or

Disclosure; Fig 1; 523pp; English.

CC any one of 1601 sequences (see ABP82019 to ABP83619) of 12-24 amino CC any one of 1601 sequences (see ABP82019 to ABP83619) of 12-24 amino CC acids. Also described: (1) an assay for the detection of a particular GC protein-coupled receptor (GPCR) or a candidate polypeptide in a sample; CC and (2) an isolated antibody having high specificity and high affinity or a cardity for a particular GPCR. (I) can be used as GPCR modulators and in CC antibody against a particular GPCR. (I) can be used as GPCR modulators and in the presence or absence of corresponding GPCRs are useful in detecting an CC antibodies. The peptides and antibodies are also useful for detecting the presence or absence of corresponding GPCRs. The antigenic peptides for CC GPCRs and antibodies are useful for dagnosing and designing drugs for CC treating immune-related diseases, growth-related diseases, cell cc diseases, or autoimmune diseases, e.g. AIDS, Alzheimer's disease, catherosclerosis, bacterial, fungal, protozoan or viral infections, catherosclerosis, bacterial, fungal, protozoan or viral infections, costeoporosis, cancer, cardiomyopathy, chronic and acute cliflammation, allergies, Crohn's disease, diabetes, graft versus host cdisease, parkinson's disease, multiple sclerosis, pain, psoriasis, cancer, depression, schizophrenia, dementia, mental retardation, memory CC hypotension, renal disorders, rheumatoid arthritis, trauma, ulcers, or cused in immunoassays and immunodiagnosis. ABZ42523 to ABZ42869 encode used in immunoassays and immunodiagnosis, ABZ42523 to ABZ42869 encode GPCR protesins given in ABP81675 to ABB82018, which are used in the exemplification of the present invention

Sequence 322

Score 1642; BB 6

RESULT 13 ABP81750

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241

RIHLDWKVLFCHVHLVSIFLSALNSSANFIIYFFVGSFRORONRONLKLVLQRALQDTPE

VDEGGGWLPQETLELSGSKLEQ 322

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RESULT 14
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Matches
The present invention relates to an isolated mas-related gene D (MrgD) polypeptide. The MrgD polypeptides are useful as therapeutics or for identifying compounds, i.e. agonists or antagonists, that alter pain perception in a mammal. The compounds are useful for treating impaired sensory perception, e.g. chronic intractable pain or neuropathic pain, promoting wound healing, restoring normal sensitivity following injury, or treating ocular conditions, particularly those associated with pressure such as glaucoma. The Mrg genes or proteins may be used as
                                                                                                                                                                  New mas-related gene D polypeptides, useful as therapeutics or in identifying agonists or antagonists that alter pain perception in a mammal for treating impaired sensory perception, e.g. chronic intractable pain or neuropathic pain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mas-related gene D;
sensory perception;
                                                                                                                                       Disclosure; SEQ ID NO 31; 220pp; English
                                                                                                                                                                                                                                                                                                                                                        26-JUN-2002;
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glaucoma; Mrg.
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Best Local Simi
Matches 313;
                                                                                     HI7T213 protein, encoded pathological mechanism, c preventives or remedies f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HI7T213; transgenic; G protein-coupled receptor; GPCR; ophthalmological; cytostatic; nephrotropic; antiinflammatory; dermatological; analgesic; vulnerary; neuroprotective; human; receptor.
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                                                                       dermatitis.
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DB; ADO44603.
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97.2%;
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Query Match 97.1%; Score 1642; DB 8; Length 322; Best Local Similarity 97.2%; Pred. No. 5.2e-170; Matches 313; Conservative 5; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a non-human mammal that carries a DNA integrated with a foreign HI7T213 or its mutant gene, or a part of it. The non-human animal is particularly a rat. Such gene shows phenotypes of e.g. cataract conset, transient skin rash and proliferation-promoting activity. The foreign HI7T213 gene is a gene that encodes a G protein-coupled receptor (GPCR) protein HI7T213. The protein, its encoded DNA and constructed transgenic animals are useful for clarifying pathological mechanism, developing therapeutic methods and screening preventives or remedies for related diseases e.g. cataract, cancer, and dermatitis. The present sequence represents a human HI7T213 protein.
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Search completed: February 3, 2006, 20:27:05 Job time : 205 secs

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Minimum
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	œ	7	σ	ហ	4	ω	2	–	No.	DAC::] +
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165989	A41783	A37963	JC2434	JN0605	A48227	S36143	JN0763	I57955	S42096	JC5835	B41795	A55733	JC2083	B42009	I57940	A53611	A53752	S29248	A45291	D41795	C42009	A23669	JQ1231	A35639	851001	TVHUAS	TVRTAS	94	ID	
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391	391	391	359	371	353	350	375	352	364	384	380	384	380	371	352
N	N	N	N	N	N	N	N	س	N	N	N	N	N	N	N
A39297	C41795	A41795	A48921	JC5796	JC2492	A39445	JC5069	\$27357	A49542	JC4629	JC2338	A47249	A55259	JC5498	A46520
somatostatin recep	somatostatin recep	somatostatin recep	interleukin-8 rece	probable chemoattr	G protein-coupled	interleukin-8 rece	G protein-coupled	complement C5a ana	N-formyl peptide c	somatostatin recep	kappa opioid recep	brain-specific som	kappa opioid recep	G protein-coupled	N-formyl peptide r

ALIGNMENTS

A39485

A39485	
Cransforming protein (mrg) - human	
C;Date: 28-Feb-1992 #sequence revision 17-Apr	#sequence revision 17-Apr-1993 #text change 09-Jul-2004
C;Accession: A39485	
R; Monnot, C.; Weber, V.; Stinnakre, J.; Bihoreau, C.; Teutsch, B.; Corvol, P.; Clauser, F.	au, C.; Teutsch, B.; Corvol, P.; Clauser, I
Mo1. Endocrino1. 5, 1477-1487, 1991 A;Title: Cloning and functional characterization of a novel mas-related gene, modulating	on of a novel mas-related gene, modulating
A;Reference number: A39485; MUID:92130997; PMID:1723144	D:1723144
A;Accession: A39485	
A;Status: preliminary	
A; Molecule type: DNA	
A; Residues: 1-378 < MON>	
A;Cross-references: UNIPROT:P35410; UNIPARC:UPI000003B44C; GB:S78653; NID:g244209; PIDN:	I000003B44C; GB:S78653; NID:g244209; PIDN:I
C:Superfamily: mas transforming protein	perbrane protein
	Part Carlo
Query Match 26.8%; Score 453; DB 2; Best Local Similarity 36.7%; Pred. No. 5.9e-30; Matches 105; Conservative 55; Mismatches 8	; Score 453; DB 2; Length 378; ; Score 453; DB 2; Length 378; 55: Mismatches 80; Indels 46: Gaps 9;
 Qy 36 IISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHI	LVAANFLFLSGHI 80
Db 84 LVSLCGVLLNGTVFWLLCCG-ATNPYMVYILHLVAADVIYLCCSAVGFLQVTLLTYHGVV 142	LVAADVIYLCCSAVGFLQVTLLTYHGVV 142
 Qy 81 IFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTERCLSILWPIWYHCRRPRYLSS	LSAISTERCLSILWPIWYHCRRPRYLSS 140
 Db 143 FFIPDFLAILSPFSFEVCLC	DFLAILSPFSFEVCLCLLVAISTERCVCVLFPIWYRCHRPKYTSN 191
 Qy 141 VMCVLLWALSILENNFCDFLFSGANSVWCBTSDFITIAWLVFLCVVLCGSS	CETSDFITIAWLVFLCVVLCGSS 195
Db 192 VVCTLIWGLPFCINIVKSLFLTYWKHVKACVIFLKLSGLFHAILSLVMCVSS	KHVKACVIFLKLSGLFHAILSLVMCVSS 243
 QY 196 LVLLVRILCGSRKMPLTRLYVTILLTVLVEFLCGLPFGIQWALFSRIHLDWKVLFCHVHL	CGLPFGIQWALFSRIHLDWKVLFCHVHL 255
 Db 244 LTLLIRFLCCSQQQKATRVYAVVQISAPMFLLWALPLSVAPLITDFKMFVTTSYL	WALPLSVAPLITDFKMFVTTSYL 298
Qy 256 VSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPEV 301	KLVLQRALQDTPEV 301
 Db 299 ISLFL-IINSSANPIIYFFVGSLRKKRLKESLRVILQRALADKPEV 343	RVILQRALADKPEV 343

RESULT TVRTAS

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A;Gene: mas
C;Superfamily: mas transforming protein
C;Superfamily: mas transforming protein
C;Keywords: G protein-coupled receptor; transforming protein; transmembrane
F;31-47/Domain: transmembrane #status predicted <TM1>
F;72-88/Domain: transmembrane #status predicted <TM2>
F;149-165/Domain: transmembrane #status predicted <TM4>
F;185-204/Domain: transmembrane #status predicted <TM4>
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A;Accession: A31816
A;Molecule type: mRNA
A;Residues: 1-324 <YOU>
A;Cross-references: UNIPROT:P12526; UNIPARC:UPI0000043DE7; GB:J03823; NID:g205313; C;Genetics:
                                                                                                F;105-135/Domain: transmembrane #status predicted <TM3>
F;150-172/Domain: transmembrane #status predicted <TM4>
F;156-214/Domain: transmembrane #status predicted <TM5>
F;186-214/Domain: transmembrane #status predicted <TM6>
F;255-250/Domain: transmembrane #status predicted <TM6>
F;258-286/Domain: transmembrane #status predicted <TM7>
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C;Superfamily: mas transforming protein
C;Keywords: G protein-coupled receptor; glycoprotein; proto-oncogene; transforming
F;31-61/Domain: transmembrane #status predicted <TM1>
F;66-97/Domain: transmembrane #status predicted <TM2>
                                                                                                                                                                                                                                                                                                                                                    A;Cross-references:
C;Genetics:
                                                                                                                                                                                                                                                                                                             A; Gene: GDB: MAS1
A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-325 < YOU>
                                                                                                                                                                                                                                                                                                                                                                                                                                     A,Title: Isolation and characterization of a new cellul A,Reference number: A01375; MUID:86218084; PMID:3708691 A,Accession: A01375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R; Young, D.; Waitches, G.; Birchmeier, Cell 45, 711-719, 1986
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Query Match
Best Local S
Matches 107
                                                                               ;225-250/Domain: transmembrane;258-286/Domain: transmembrane;5,16,22,272/Binding site: carb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Species: Homo sapiens (man)
;Date: 04-Dec-1986 #sequence_revision 04-Dec-1986 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SSANPFIYFFVGSSKKKRFRESLKVVLTRAFKDEMQPRRQEGNG
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                     25.4%;
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                                                                                 carbohydrate
  63;
  Score 429.5;
Pred. No. 4.4e
53; Mismatches
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                                                                                   (Asn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a new cellular oncogene
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                       4.4e-28
                                                                                 (covalent)
                                          DB 1; Length 325;
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                                                                                                                                                                                                                                                                                                                         SIYILNLVAANELELSGHIIES-----PLPLINIRHPISKILSPVMTFPYFIGLSMLSAI
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                                                                                               VLQRALQD 297
                                                                                                                           MPMRLLYLLYYEY---
                                                                                                                                                  LPFGIQWALESRIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKL
                                                                                                                                                                                 DCRAVIIFIAILSFLVFTPLMLV-SSTILVVKIRKNTWASHSSKLYIVIMVTIIIFLIFA
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                                                                       VLTRAFKD 305
  mouse
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A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-87,'I',89-324 <RES> A;Cross-references: UNIPARC:UPI0000029856; EMBL:X67735; C;Genetics: A;Gene: mas C;Superfamily: mas transforming protein C;Keywords: G protein-coupled receptor; proto-oncogene; Nilternate names: mas proto-oncogene protein; probable G protein-coupled receptor C;Species: Mus musculus (house mouse)
C;Date: 10-Apr-1996 #sequence revision 19-Apr-1996 #text_change 09-Jul-2004
C;Accession: S51001; 148647; S29619
R;Metzger; R.; Bader, M.; Ludwig, T.; Berberich, C.; Bunnemann, B.; Ganten, D.
FEBS Lett. 357, 27-32, 1995
A;Title: Expression of the mouse and rat mas proto-oncogene in the brain and periph A;Reference number: S51001; MUID:95094925; PMID:8001672
A;Accession: S51001 R;Metzger, R.; Bader, M.; Ludwig, T.; Berberich, C.; Bunneme FBBS Lett. 357, 27-32, 1995
A;Title: Expression of the mouse and rat mas proto-oncogene A;Reference number: I48647; MUID:95094925; PMID:8001672
A;Accession: I48647 A,Status: nucleic acid sequence not shown A,Molecule type: DNA A;Cross-references: UNIPROT:P30554; UNIPARC:UPI000003B44B; EMBL:X67735 R;Metzger, R.; Bader, M.; Ludwig, T.; Berberich, C.; Bunnemann, B.; Ganten, A;Residues: Matches Query Match Local 1-324 <MET> Similarity Conservative 25.0%; Score 422.5; DB 2; 37.7%; Pred. No. 1.7e-27; 55 Mismatches proto-oncogene; transmembrane 99; Length Indels NID:g53011; ä 324; the brain and peripheral t 23; brain and peripheral PIDN:CAA47964.1; Gaps protein Ö PID

205 161

GSRKMPLTRLYVTILLTVLVFLLCGLPFGIQMALFSRIHLDWKVLFCHVHLVSIFLSALN : | : | : | : | : | : | : | : | : |

101

92 41 37

HPISKILSPVMTFPYFIGLSMLSAISTERCLSILWPIWYHCRRPRYLSSVMCVLLWALSL | SPLGFVENGILLWFLCFRMRRNPFTVYITHLSMADISLLFCIFILSTDYALDYELSSGH 100 ISLVALTGNAVVLMLLGCRMRRNAVSIYILNLVAANFLFLSGHIJFS-----PLPLINIR

151

LRSILEWMFCDFLFSGANSVWCETSD-----FITI-AWLVFLCVVLCGSSLVLLVRILC HYTIVTLSVTFLFGYNTGLYLLTAISVERCLSVLYFIWYTSHRPKHQSAFVCALLCALSC 160

LVTTMEYVMC--IDSGEES--HSRSDCRAVIIFIAILSFLVFTPLMLVSSS-ILVVKIRK

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A35639
G protein-coupled rec
C; Species: Rattus nor
C; Date: 22-Jan-1993 †
C; Accession: A35639
R; Ross, P.C.; Figler.
                    A; Molecule type: DNA
A; Residues: 1-355 <BEC>
A; Cross-references: UNIPROT: P21109; UNIPARC: UPI000012D4ED;
A; Cross-references: UNIPROT: P21109; UNIPARC: UPI000012D4ED;
A; Lee, J.; Kuang, W.J.; Rice, G.C.; Wood, W.I.
J. Immunol. 148, 1261-1264, 1992
A; Title: Characterization of complementary DNA clones encod A; Reference number: A46483; MUID: 92148149; PMID: 1737938
A; Accession: A46483
A; Accession: A46483
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-355 <LEE>
A; Residues: 1-355 <LEE
A; Residues
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A;Title: RTA, a candidate G protein-coupled receptor: cloning, sequencing, and tiss A;Reference number: A35639; MUID:90222168; PMID:2109324
A;Accession: A35639
A;Molecule type: mRNA
A;Residues: 1-343 <ROS>
A;Cross-references: UNIPROT:P23749; UNIPARC:UPI0000046F10; GB:M35297; NID:g206809; C;Superfamily: mas transforming protein
C;Keywords: G protein-coupled receptor; glycoprotein; membrane protein
C;Keywords: G protein-coupled receptor; glycoprotein; membrane protein
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R;Beckmann, M.P.; Munger, W.E.; Kozlosky, C.; VandenBos, T.; Price Biochem. Biophys. Res. Commun. 179, 784-789, 1991
A;Title: Molecular characterization of the interleukin-8 receptor A;Reference number: JQ1231; MUID:91378994; PMID:1898400
A;Accession: JQ1231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin-8 receptor - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 31-Mar-1992 #sequence revision 31-Mar-1992
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    A;Cross-references:
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Species: Rattus norvegicus (Norway rat)
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    UNIPARC: UPI000012D4ED;
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                                                                                                                                                                                               DNA clones encoding PMID:1737938
GB:M82873; NID:g165440; PIDN:AAA31376.1;
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R;fhomas, K.M.; Pyun, H.Y.; Navarro, J.
J. Biol. Chem. 265, 20061-20064, 1990
A;Title: Molecular cloning of the fMet-Leu-Phe receptor from neutrophils.
A;Reference number: A23669; MUID:91056034; PMID:1700779
A;Accession: A23669
A;Accession: A2369
A;Accession: A2369
A;Accession: A2369
A;Accession: Cype: mRNA
A;Residues: 1-354 <-YHO>
A;Cross-references: UNIPROT:P21109; UNIPARC:UPI0000156FB7; GB:M58021; GB:J05705; NID:g165
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; glycoprotein; membrane protein; neutrophil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interleukin-8 receptor, high affinity - rabbit N;Alternate names: fMLP receptor C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 C;Accession: A23669
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A23669
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A;Note: sequence extracted from NCBI backbone (NCBIN:81526, NCBIP:81530)
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; transmembrane protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                118 TERCLSILWPIWYHCRR----PRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANS--V
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                                                                                                                                                                                                                                           64 YILNLVAANFLFLSGHIIFSPL--PLINIRHPISKILSPVMTFPYFIGLSMLSAISTERC 121
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                                                                                                                                                                                                                                                                                  23 TGMPPVE-KDYSPCLVVTQTLNKYVVVVIYALVFLLSLLGNSLVMLVILYSRSNRSVTDV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          82 YLLNLAMADLLF----ALTMPIWAVSKEKGWIFGTPLCKVVSLVKEVNFYSGILLLACIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64 YILNLVAANFLFLSGHIIFSPLPLIN-----IRHPISKILSPVMTFPYFIGLSMLSAIS
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                                                                                                                                                                                                                                                                                                                                                                96;
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                                                                                                                                                            LSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANS--VWCETSDFI 179
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    FLLCGLPFG----
                                                                               TIAWLVFLCVVL-----CGSSLVLLVRILC-GSRKMPLTRLYV-----
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                                         TAKW----CMVLRILPHTFGFILPLLVMLFCYGFTLRTLFQAHMGQKHRAMRVIFAVVLI
                                                                                                                      LAIVOST-RTLTOKRHLVKFICLGIWALSLILSLPFFLFRQ-VFSPNNSSPVCYEDLGHN 198
                                                                                                                                                                                                   YLLNLAMAP-AFCPDHAYLGRLQGKRLDFRTPLCKVVSLVKEVNFYSGILLLACISVDRY 140
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Pred. No. 3.7e-12;
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    IQWALFSRIHLDWKVLFCHVHLVSIFLSALNSSANPII
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somatostatin receptor 2 - mouse C;Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Date: 30-Sep-1993 #sequence revision 30-Sep-1993 #text_change 09-Jul-2004 C;Accession: D41795; I56236 R;Yamada, Y; Post, S.R.; Wang, K.; Tager, H.S.; Bell, G.I.; Seino, S. Rroc. Natl. Acad. Sci. U.S.A. 89, 251-255, 1992 A;Title: Cloning and functional characterization of a family of human and mouse A;Reference number: A41795; MUID:92108031; PMID:1346068 A;Accession: D41795
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C; Superfami
C; Keywords:
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C;Comment: This fMet-Leu-Phe receptor homolog, whose ligand is not yet known, appears
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R.Bao, L.; Gerard, N.P.; Eddy Jr., R.L.; Shows, T.B.; Genomics 13, 437-440, 1992
A;Title: Mapping of genes for the human C5a receptor (C: A;Reference number: A42009; MUID:92307681; PMID:1612600
A;Accession: C42009
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A;Map position: 19q13.3-19q13.4
A;Introns: #status absent
C;Superfamily: vertebrate rhodopsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Gene: GDB:FPRL2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-353 <BAO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N;Alternate
C;Species: F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Species: Homo sapiens (man)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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Pred. No. 1.9e-11;
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A;Status: nucleic acid sequence not shown
A;Molecule type: DNA
A;Residues: 1-369 <YAM>
A;Residues: 1-369 <YAM>
A;Cross-references: UNIPROT:P30875; UNIPARC:UPI0000000A47; GB:M81832; NID:g201060;
A;Cross-references: UNIPROT:P30875; UNIPARC:UPI000000A47; GB:M81832; UNIPARC:UPI000000A47; GB:M81832; UNIPARC:UPI000000A47; GB:M81832; UNIPARC:UPI000000A47; GB:M81832; UNIPARC:UPI00000A47; GB:M81832; UNIPARC:UPI00000A47; GB:M81832; UNIPARC:UPI00000A47; GB:M81832; UNIPARC:UPI00000A47; GB:M81832; UNIPARC:UPI00000A47; GB:M81832; UNIPARC:UPI0000A47; GB:M81832; UNIPARC:UPI000A47; GB:M81832; UNIPARC:UPI000A47; GB:M81832; UNIPARC:UPI000A47; UNIPARC:UPI000A47; UNIPARC:UPI000A47; UNIPARC:UPI000A47; UNIPARC:UPI000A47; UNIPARC:UPI000A47; UNIPARC:UPI000A47; UNIPAR
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A;Title: T lymphocytes isolated from the hepatic granulomas of schistosome-infected A;Reference number: I56236; MUID:94300079; PMID:7913111
A;Accession: I56236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Expression cloning of a rat brain sc
A;Reference number: A45291; MUID:92262491; PM
A;Accession: A45291
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A;Residues: 99-309 <RES>
A;Cross-references: UNIPARC:UPI00001778F0; GB:S71756; NID:g560631
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; hormone receptor; transme
                                                                                                                                                                                                                                                                                                                                               A;Residues: 1-369 <KLU>
A;Cross-references: UNIPROT:P30680; UNIPARC:UPI0000135FF7; GB:M93273; NID:g207026; PIDN:IA;NOTe: sequence extracted from NCBI backbone (NCBIN:102315, NCBIP:102316)
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     somatostatin receptor, somatotropin release-inhibiting factor receptor, SRIF receptor C;Species: Rattus norvegicus (Norway rat) C;Date: 25-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004 C;Accession: A45291 C;Accession: A45291 R;Kluxen, F.W.; Bruns, C.; Lubbert, H.
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Best Local &
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Natl. Acad. Sci. U.S.A. 89, 4618
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Similarity 28.0%; Pred. No. 5.5e-11;
94; Conservative 58; Mismatches 129;
                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                     LGTKLTPINGREET-PCYNQTLS--FTGLTCIISLVALTGNAVVLW-LLGCRMRRNAVSI
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   YILNLVAANFLFLSGHIIFSPLPLINIRHPISK-ILSPVMT---FPYFIGLSMLSAISTE
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PMID:1374909
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isotype E.; Navarro

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A; Molecule type: mRNA
A; Residues: 1-346 < VAN's
A; Residues: 1-346 < VAN's
A; Cross-references: UNIPROT: P30875; UNIPARC: UPI0000044D82; EMI
C; Superfamily: vertebrate rhodopsin
C; Keywords: G protein-coupled receptor; transmembrane protein
                                                                       RESULT
A53752
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S29248
interleukin-8 receptor (clone 5Bla) - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: A53752
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A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R; Vanetti, M.; Kouba, M.; Wang, X.; Vogt, G.; Hoellt, FEBS Lett. 311, 290-294, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      somatostatin receptor 2B - mouse
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
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28.0%;
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Pred. No. 1.:
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A;Molecule type: mRNA
A;Residues: 1-15 <RE2>
A;Cross-references: UNIPARC:UPI00000053D; EMBL:U11872; NID:g511808; PIDN:AAA64380.1;
A;Cross-references: UNIPARC:UPI000000053D; EMBL:U11877; NID:g511818; PID:
14; PID:g511815; EMBL:U11876; NID:g511816; PID:g511817; EMBL:U11877; NID:g511818; PID:
R.Sprenger, H.; Lloyd, A.R.; Lautens, L.L.; Bonner, T.I.; Kelvin, D.J.
J. Biol. Chem. 269, 11065-11072, 1994
J. Biol. Chem. 269, 11065-11072, 1994
A;Fitle: Structure, genomic organization, and expression of the human interleukin-8 relapidation: A53611; MUID:94209273; PMID:7512557
A;Accession: A53611
A;Status: preliminary
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J. Biol. Chem. 269, 12391-12394, 1994
A;Title: Molecular characterization of a novel rabbit interleukin-8 receptor isotyp
A;Reference number: A53752; MUID:94230294; PMID:8175642
A;Accession: A53752
A;Status: preliminary
A;Molecula type: mRNA
A;Residues: 1-358 <PRA>
A;Residues: 1-358 <PRA>
A;Residues: 1-358 <PRA>
A;Residues: 1-358 <PRA>
A;Residues: Gerences: UNIPROT:935344; UNIPARC:UPI000012D4F4; GB:L24445; NID:g437661;
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        interleukin-8 receptor type B - human (75pecies: Homo sapiens (man) (75pecies: Homo sapiens (man) (75pecies: Homo sapiens (75pecies: Homo sapiens (75pecies: 1370) 4 #sequence revision 12-Apr-1996 #text_change 09-Jul-2004 (75pecies: 137898; 138712; \bar{A}53611; \bar{A}39446 R; \bar{A}446 R; \bar{A}5.K.; Shetty, A.; Tiffany, H.L.; Murphy, P.M. (75pecies: 15pecies: 137898; MUID:95014476; PMID:7929358 R.Reception: 137898; MUID:95014476; PMID:7929358
                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-360 <RES>
A;Cross-references: UNIPARC:UPI00000746D6; R;Murphy, P.M.; Tiffany, H.L. Science 253, 1280-1283, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
A53611
                                                                     A;Residues:
                                                                                                  A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary
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                                                                     6-360 <SPR>
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Pred. No. 1.7e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches 105;
                                              GB:M99412; GB:L19593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
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N;Alternate names: somatotropin release-inhibiting factor (;Species: Rattus norvegicus (Norway rat) (C;Date: 02-Aug-1996 #sequence revision 02-Aug-1996 #text_C;Accession: 157940; 157940; $39244 R;O'Carroll, A.M.; Lolait, S.J.; Konig, M.; Mahan, L.C. Mol. Pharmacol. 42, 939-946, 1992 A;Title: Molecular cloning and expression of a pituitary shareference number: 157940; MUID:93125499; PMID:1362243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: Cloning of complementary DNA encoding a functional human A;Reference number: A39446; MUID:91368200; PMID:1891716 A;Accession: A39446
                                                                                                                                          A;Cross-references: UNIPROT:P30938; UNIPARC:UPI0000135FFB;
R;O'Carroll, A.M.; Lolait, S.J.; Konig, M.; Mahan, L.C.
Mol. Pharmacol. 44, 1278, 1993
A;Title: Molecular cloning and expression of a pituitary su
A;Reference number: 157949; MUID:94088493; PMID:8264565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPARC:UPI00000746D6; GB:C;Comment: This receptor, unlike ILBRA, binds C;Genetics:
A;Genet:GDB:IL8RB; IL8RA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Status: preliminary; nucleic acid sequence not A;Molecule type: mRNA A;Residues: 6-360 <MUR>
A; Description:
                  A; Experimental source: pituitary R; Penetta, R.; Greenwood, M.; Patel, Y.C. submitted to the EMBL Data Library, Augus
                                                                                  A;Status: preliminary; translated A;Molecule type: mRNA A;Residues: 341-363 <OCA2>
                                                                                                                              A; Accession: I57949
                                                                                                                                                                                                                          A; Molecule type: mRNA
A; Residues: 1-363 < OCA1>
                                                                                                                                                                                                                                                        A; Status: preliminary; translated from GB/EMBL/DDBJ
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A; Map position: 2q35
                                                                                                                                                                                                                                                                                                                                                                                                               somatostatin receptor 5 - rat
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                                                                 Cross-references: UNIPARC:UPI00000004EB; GB:S67370; NID:g455947; PIDN:AAB29371
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DFITIAWLVFLCVV--LCGSSLVLLVRILC-GSRKMPLTRLYV-----TILLTVLVF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LAIV----HATRTLTQKRYLVKFICLSIWGLSLLLALPVLLFRRTVYSSNVSPACYEDM
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Pred.
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FMLP-related receptor 1 - human
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A; Reference number: S39244
A; Accession: S39244
A; Molecule type: mRNA
A; Residues: 309-363 <PEN>
A; Cross-references: UNIPARC: UPI00001707D8;
C; Genetics:
A; Gene: SSTR5
C; Superfamily: vertebrate rhodopsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 TPCYNQTLSFTG------LTCIISLVALTGNAVVLW-LLGCRMR
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ELSG 317
                                                                                                                                                                                                                                                                                                                                   TSIFCLMVMSVDRYLAVVHPLRSARW----RRPR-VAKMASAAVWVFSLLMSLPLLVFADV
                                                                                                                                                                                                                                                                                                                                                                                IGLSMLSAISTERCLSILWPI----WYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNAVSIYILNLVAANFLFLSGHIIFSPLPLINIRHPI----SKILSPVMTFP---YF
                                                                                         NSSANPIIYFFVG-SFRORONRONLKLVLOR--ALQDT----PEVDEGG----GWLPQETL
                                                                                                                                           RRRSEPKVTRMVVVV----VLVFVGCWLPFFIVNIVNLAFTLPEEPTSAGLYFFVVVLSYA
                                                                                                                                                                                                                                                                                    L--FSGANSVWCETSDFITIAWLVFLCV-----VLCGSSLVLLVRILCGSRKM----
                                                                                                                                                                                                                                                                                                                                                                                                                               KTVTNVÝ ILNLAVÁDVLÉMLG-----LÉFLATQNAVVSYWEFGSFLCRLVMTLDGINQF
                                            NSCANPILYGFLSDNFRQSFRK---VLCLRRGYGMEDADAIEPRPDKSGRPQATLPTRSC 352
                                                                                                                                                                                                                                       QEGWGTCNLSWPEPVGLWGAAFITYTSVLGFFGPLLVICLCYLLIVVKVKAAGMRVGSSR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TPSWNASAASSGNHNWSLVGSASPMGARAVLVPVLYLLVCTVGLSGNTLVIYVVLRHAKM
                                                                                                                                                                                        ----PLTRLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDWKVLFCHVHLVSIFLSAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12.7%; Score 215.5;
24.5%; Pred. No. 2.1
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Gene 118, 303-304, 1992

A;Title: Cloning of a cDNA encoding a receptor related to the formyl peptide ;
A;Reference number: JC1258; MUID:92380523; PMID:1511907

A;Recession: JC1258

A;Status: nucleic acid sequence not shown

A;Molecule type: mRNA

A;Molecule type: mRNA

A;Residues: 1-351 < PER>
A;Cross-references: UNIDARC:UPI0000012D0; EMBL:X63819; NID:g31460; PID:g31461

A;Experimental source: bone marrow mRNA

A;Experimental source: bone marrow mRNA

A;Fe, R. D; Cavanagh, S. L.; Quehenberger, O.; Prossnitz, E.R.; Cochrane, C.G.

Biochem. Biophys. Res. Commun. 184, 582-589, 1992

A;Title: Isolation of a cDNA that encodes a novel granulocyte N-formyl peptide
A;Reference number: JQ1521; MUID:92246937; PMID:1374236 Génomics 13, 437-440, 1992
A;Title: Mapping of genes for the human C5a receptor (C5AR), human FW A;Title: Mapping of genes for the human C5a receptor (C5AR), human FW A;Reference number: A42009; MUID:92307681; PMID:1612600
A;Accession: B42009
A;Status: nucleic acid sequence not shown
A;Reslatus: 1-263,'A',265-338,'C',340-351 <BAO>
A;Reslatus: 1-263,'A',265-338,'C',340-351 <BAO>
A;Cross-references: UNIPROT:P25090; UNIPARC:UPI00001778D8; GB:M76672
A;Note: authors translated the codons GTG for residue 15 as G1u, TCT
R;Perez, H.D.; Holmes, R.; Kelly, E.; McClary, J.; Andrews, W.H.
Gene 118, 303-304, 1992 N;Alternate names: FMLP receptor homolog FPR2; formyl peptide receptor like-1; probable C;Species: Homo sapiens (man) C;Date: 30-Sep-193 #sequence revision 14-Jul-1995 #text change 09-Jul-2004 C;Accession: B42009; JC1258; JQ1521; A42492; I54751; S21581 C;Accession: B42009; JC1258; JQ1521; A42492; I54751; S21581 C;Accession: B42009; JC1258; JQ1521; A42492; I54751; S21581 **EMLP** PID: 931461 for receptor (FPR), 19 as 얁 크

peptide

receptor

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A;Cross-references: GDB:127554; OMIM:136538
A;Map position: 19q13.3-19q13.4
A;Introns: #status absent
C;Superfamily: vertebrate rhodopsin
C;Keywords: chemotaxis; G protein-coupled receptor; glycoprotein; transmembrane protein
C;C-53/Domain: transmembrane #status predicted <TM2>
F;59-83/Domain: transmembrane #status predicted <TM3>
F;100-121/Domain: transmembrane #status predicted <TM4>
F;145-169/Domain: transmembrane #status predicted <TM4>
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A;Cross-references: UNIPARC:UPI00000012D0; GB:D10922; NID:g219864; PIDN:BAA01720.1;
A;Cross-references: UNIPARC:UPI00000012D0; GB:D10922; NID:g219864; PIDN:BAA01720.1;
C;Comment: This G-protein coupled receptor, homologous to the N-formyl peptide receptor; comment: This G-protein coupled receptor, homologous to the N-formyl peptide receptor Grand State of the N-formyl peptide receptor for some other l
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R;Nomura, H.; Nielsen, B.W.; Matsushima, K.
Int. Immunol. 5, 1239-1249, 1993
Int. Immunol. 5, 1239-1249, 1993
A;Title: Molecular cloning of cDNAs encoding a LD78 receptor and putative leukocyte chem A;Reference number: I54751; MUID:94092629; PMID:7505609
A;Accession: I54751
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A;Note: formyl peptide stimulated calcium mobilization comparable to that of the formyl
R;Murphy, P.M.; Ozcelik, T.; Kenney, R.T.; Tiffany, H.L.; McDermott, D.; Francke, U.
J. Biol. Chem. 267, 7637-7643, 1992
A;Title: A structural homologue of the N-formyl peptide receptor. Characterization and A;Reference number: A42492; MUID:92218423; PMID:1373134
A;Accession: A42492
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A; Residues: 1-351 < MUR>
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Best Local S
Matches 88
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   NPTSSLAFFNSCLNPMLYVFVGQDFRERL-IHSLPTSLERALSE
                                                                                                                                                                                                                                                                   FNFAS----WGGTPEERLKVAITMLTARGIIRFVIGFSLPMSIVAICYGLIAAKIHKKGM 233
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                                                           ----LSALNSSANPITYFFVG-SFRORQNRQNLKLVLQRALQD 297
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Search completed: February

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2006, 20:32:10

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Job time : 44 secs

iuis bade Blauk (nsbto)

Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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Minimum DB
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1: uniprot_sprot:*
2: uniprot_trembl:*
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1 MDPTIPVLGTKLTPINGREE......EGGGWLPQETLELSGSKLEQ 322
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"Generation and initial analysis of more than 15,000 full-length human	S.J.M., Marra M.A.;	Butterfield Y.S.N. Krzywinski M.I., Skalska U., Smailus D.E.,	Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,	Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,	Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,	Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,	Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,	Romak S.A. McGwan D.J. McKernan K.J. Malek J.A. Gunarathe P.H	Brownstein M.J., Usdin T.B., Tosniyuki S., Carninci P., Frange C.,	Bonaldo M.F., Casavant T.L., Scheet	, Marusina K., Farmer A.A., Rubin G.M., Hong L.,	Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,	Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,	Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,	Strainshard R I. Feingold R A. Groise I. H. Derge I. G.	######################################	NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].		Nat. Neurosci. 5:201-209(2002).	•	"Proenkephalin A gene products activate a new family of sensory	ker P., Ahmad S.;	lin M., Fortin Y., Banville D., Shen S., Stroem P., Payza	J. Hoffert C. Cao J. Schmidt R. Pelletier M.	O'Donnell D. Rov MO.	MRDI.TMR=21843733 - Ph/Med=11850634 - DOT =10 1038/mal5:	NICT FOR THE CROTTENCE VARIANT AGN-160 AND TISSIE CRECIPICITY	Ceil 106:619-632(2001).	nociceptive sensory neurons.";	iverse family of GPCRs expressed in specific subsets	mon M.I., Anderson	MEDLINE=21435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4;	NICTEOTIDE SECTIONCE	NCBL TAXLD=9606;	Homo	lia; Eutheria; Euarchontoglires; Primates; Cata	ryota; Mo	Homo sapiens (Human).	Name=MRGPRX3: Synonyms=MRGX3. SNSR1:		48, Last annotation update)	. 45, Last	45, Created)			RESULT 1

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PRINTS; PR00237; GFCRHODOPSN.

PROSITE; PS00237; G PROTEIN RECEP_F1_1; 1.

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.

G-protein_coupled_receptor; Polymorphism; Receptor; Transducer;
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EMBL; AF474987; AAL86878.2; -; Genomic_DNA.
EMBL; BC067292; AAH67292-1; -; mRNA.
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FUNCTION: Orphan receptor. Probably involved in the function nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Potently activated by enkephalins (By similarity).

SUBCELLULAR LOCATION: Integral membrane protein.

TISSUE SPECIFICITY: Uniquely localized in a subset of small root and trigeninal sensory neurons.

SIMILARITY: Belongs to the G-protein coupled receptor 1 fami
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Cytoplasmic (Potential).

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Cytoplasmic (Potential).

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Extracellular (Potential).

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Cytoplasmic (Potential).

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/FTId=VAR 019434.

S -> P (in Ref. 2).

C -> R (in Ref. 3).

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R -> K (in Ref. 2).
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Q8TDE0;
25-OCT-2004
25-OCT-2004
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MEDLINE=21853733; PubMed=11850634; DOI=10.1038/nn815;

Lembo P.M.C., Grazzini E., Groblewski T., O'Donnell D., I

Zhang J., Hoffert C., Cao J., Schmidt R., Pelletier M., I

Gosselin M., Fortin Y., Banville D., Shen S., Stroem P.,

Dray A., Walker P., Ahmad S.;

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PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Glycoprotein; Receptor; Transducer;
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Ensembl; ENSG00000179826; Homo sapiens
InterPro; IPR000276; GPCR_Rhodpsn.
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Eukaryota; Metazoa;
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Sensory neuron-specific
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Neurosci. 5:201-209 (2002).

PUNCTION: Orphan receptor. Probably involved in the function routiceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Potently activated by enkephalins (By similarity).

SUBCELLULAR LOCATION: Integral membrane protein.

TISSUE SPECIFICITY: Uniquely localized in a subset of small
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                                                                                                                                                                                                                                                                                                                                     Swiss-Prot entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no restr
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                                                                                                                                      The process of the products activate a new family of sensory of neuron-specific GPCRs."; (2002).

IN Neurosci. 5:201-209 (2002).

IN Neurosci. 5:201-209 (2002).

Concidentive neurons. May regulate nocidentor function of nocidentive neurons. May regulate nocidentor function and/or development, including the sensation or modulation of pain.

Concording activated by enkephalins including BAM22 (bovine adrenal medulla peptide 22) and BAM (8-22). BAM22 is the most potent compound and evoked a large and dose-dependent release of intracellular calcium in stably transfected cells. G(alpha) que contents are involved in the calcium-signaling pathway.

Contents SPECIFICITY: Uniquely localized in a subset of small dorsal contents.
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CARBOHYD
SEQUENCE
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Zhang J., Hoffert C., Cao J., Schmidt R., Pelletler M., Labarre
Gosselin M., Fortin Y., Banville D., Shen S., Stroem P., Payza I
Dray A., Walker P., Ahmad S.;
                            This
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MEDLINE=21853733; PubMed=11850634; DOI=10.1038/nn815;
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  copyright. It is produ
tute of Bioinformatics
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N-linked (GlcNAc. . .) (P
D8C24308E3B4611B CRC64;
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Pred. No. 9
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rimates; Catarrhini; Hominidae;
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Best Local S
Matches 267
                   MRGX1 HUMAN STANDARD; PRT; 322 AA.

Q96LB2; Q8TDD8;
25-OCT-2004 (Rel. 45, Created)
25-OCT-2004 (Rel. 45, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
Mas-related G-protein coupled receptor member X1
specific G-protein coupled receptor 4).
Name=MRGPRXI; Synonyms=MRGX1, SNSR4;
Homo sapiens (Human).
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PROSITE; PS00237; G PROTEIN_RECEP_F1_2; 1.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Glycoprotein;
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Ensembl; ENSG00000170255; Homo sar
InterPro; IPR000276; GPCR_Rhodpsn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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4 (Potential).

Extracellular (Potential).

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Cytoplasmic (Potential).

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Extracellular (Potential).

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Nolinked (GlCMAC. . .) (Potential).
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1 (Potential).
Cytoplasmic (Potential).
2 (Potential).
Extracellular (Potential).
3 (Potential).
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Pred. No. 1.9e-92;
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                                                         (Sensory
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Eukaryota; Metazoa;

Eutheria;

Chordata; Craniata; Vertebra Euarchontoglires; Primates;

Vertebrata;

Catarrhini; Hominidae;

Euteleostomi;

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G-protein co
Transducer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE, AND TISSUE SPECIFICITY.
MEDLINE=21853733; PubMed=11850634; DOI=10.1038/nn815;
Lembo P.M.C., Grazzini E., Groblewski T., O'Donnell D., I
Zhang J., Hoffert C., Cao J., Schmidt R., Pelletier M., I
Gosselin M., Forrin Y., Banville D., Shen S., Stroem P.,
Dray A., Walker P., Ahmad S.;
                                                                                                                                                                                                                                                                                                                                         the
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Cell 106:619-632(2001).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=22040266; PubMed=12044878; DOI=10.1016/S0014-5793(02)02775-8; Takeda S., Kadowaki S., Haga T., Takaesu H., Mitaku S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dong X., Han S.-K., Zylka M.J., Simon M.I., Ander "A diverse family of GPCRs expressed in specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo.
NCBI_TaxID=9606;
                                                                                                    Pfam; PF00001; 7tm 1; 1.

PRINTS; PR00237; GFCRHODDPSN.
PROSITE; PS00237; G PROTEIN_RE
PROSITE; PS50262; G PROTEIN_RE
G-protein coupled receptor; Gl
                                                                                                                                                                                                      Ensembl; ENSG00000170255; HGNC; HGNC:17962; MRGPRX1.
                                                                                                                                                                                                                                     EMBL; AY042213; AAK91804.1; -; EMBL; A4747490; AAL86881.1; -; EMBL; AB083628; BAB89341.1; -; EMBL; AB065846; BAC06064.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        receptor genes."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Suwa M., Sato T., Okouchi I., Arita M., Futami K., Matsumoto
Tsutsumi S., Aburatani H., Asai K., Akiyama Y.;
"Genome-wide discovery and analysis of human seven transmemb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Proenkephalin A gene products activate a neuron-specific GPCRs.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=21435808;
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                                                                                                                                                                           nterPro; IPR000276; GPCR_Rhodpsn.
                                                                                                                                                                                                                                                                                                                                                                                                                nociceptive neurons. May regulate nociceptor function of nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Potently activated by enkephalins including BAM22 (bovine adrenal medulla peptide 22) and BAM (8-22). BAM22 is the most potent compound and evoked a large and dose-dependent release of intracellular calcium in stably transfected cells. G(alpha)q proteins are involved in the calcium-signaling pathway.

TISSUE SPECIFICITY: Uniquely localized in a subset of small dorsal root and trigeninal sensory neurons.
                                                                                                                                                                                                                                                                                                                                          European
                                                                                                                                                                                                                                                                                                                                                                                                    Mas subfamily.
                                                                                                                                                                                                                                                                                                                     Swiss-Prot entry is copyright. It is proude the Swiss Institute of Bioinformatics een the Swiss Institute of Bioinformatics Institute. There a European Bioinformatics in no way modific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lett.
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435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4;
435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4;
435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4;
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 Transmembrane.
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                                                                                                                                                                                                                       Homo
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_RECEP_F1_2; 1.
_Glycoprotein; Polymorphism;
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                                                                                                                                                                         Lembo P.M.C., Grazzini E., Groblewski T., O'Donnell D., Zhang J., Hoffert C., Cao J., Schmidt R., Pelletier M., Gosselin M., Fortin Y., Banville D., Shen S., Stroem P., Paya A., Walker P., Ahmad S.;
"Proenkephain A Gena T., Price Species of the property 
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25-OCT-2004 (Rel. 45, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Sensory neuron-specific G-protein coupled rec
Nat. Neurosci. 5:201-209(2002).

-i-FUNCTION: Orphan receptor. Probably involved in the nociceptive neurons. May regulate nociceptor functi development, including the sensation or modulation potently activated by enkephalins (By similarity).

-i- SUBCELLULAR LOCATION: Integral membrane protein.
                                                                                                                                "Proenkephalin A gene products activate neuron-specific GPCRs.";
Nat. Neurosci. 5:201-209(2002).
                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9606;
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PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Glycoprotein; Receptor; Transducer;
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Ensembl; ENSG00000179817; Homo sapiens.
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83.1%;
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5 (Potential).
Cytoplasmic (Potential).
6 (Potential).
Extracellular (Potential).
7 (Potential).
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Cytoplasmic (Potential).
2 (Potential)
Extracellular (Potential).
3 (Potential).
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N-linked (GlcNAc...) (Potential).
3D6FFB4B5DDFDD90 CRC64;
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Pred. No. 3
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  sequence update)
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3.2e-90;
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81.7%; Pre
ative 23;
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RA Strausberg R.D., Collins F.S., Wagner L. B., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Rodrigues S., Sanchez A.,
RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Mihting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Moriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
ROGROSTIC A., Schein J.E., Jones S.J.M., Marra M.A.;
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PRINTS; PR00237; GPCRRHODOPSN.

PROSITE; PS00237; GPCOTEIN_RECEP_F1_1; UNKNOWN_1.

PROSITE; PS00262; G_PROTEIN_RECEP_F1_2; 1.

G-protein coupled receptor; Receptor; Transducer; SEQUENCE 322 AA; 36236 MW; C7F219F4418E8AD1 CR
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Name=MRGPRX1;
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.
-!- SUBCELLULAR LOCATION: Integral membrane protein (By EMBL, BC096835; AAH96835.1; -; mRNA.
Interpro; IPRO101276; GPCR_Rhodpsn.
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
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                                   RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                                                                                                                                                                                                                                                           CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
                                                                                                                                                                                                                                                                                                                                                                                                               FSIYILNLAAADFLFLSGRLIYSLLSFISIPHTISKILYPVMMFSYFAGLSFLSAVSTER
                                                                                                                                                                                                                                                                                                                                                                                                                                               VSIYILNLVAANFLFLSGHIIFSBLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
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WIHVDREVLECHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDASE
                                                                                                                                                                    IAMLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
                                                                                                                                                                                                                                                                        CLSVLWPIWYRCHRPTHLSAVVCVLLWALSLLRSILEWMLCGFLFSGADSAWCQTSDFIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MDPTISTLDTELTPINGTEETLCYKQTLSLTVLTCIVSLVGLTGNAVVLMLLGCRWRRNA
                                                                                                                                    VAWLIFLCVVLCGSSLVLLIRILCGSRKIFLTRLYVTILLTVLVFLLCGLFFGIQFFLFL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 1370; DI
Pred. No. 3.2e
23; Mismatches
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VDEGGGQLPEEILELSGSRLDQ

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RX MEDLINE=2238257; PubMed=12477922; DOI=10.1073/pnas.242603899;
RX MEDLINE=2238257; PubMed=12477922; DOI=10.1073/pnas.242603899;
RA Strausberg R.D., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L.H., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L.H., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Distchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Distchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
"Generation and initial analysis of more than M.A.;
"Generation and initial analysis of more than 15,000 full-length human full modes cDNA sequences.";

Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El the European Bioinformatics Institute. There are no restruse as long as its content is in no way modified and this use as long as its content is in no way modified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  [3]
NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Proenkephalin A gene products activate neuron-specific GPCRs.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lembo P.M.C., Grazzini B., Gr
Zhang J., Hoffert C., Cao J.,
Gosselin M., Fortin Y., Banvi
Dray A., Walker P., Ahmad S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE [GENOMIC DNA], MEDLINE=21853733; PubMed=11850634;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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"A diverse family of GPCRs expressed in specific subsets of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [GENOMIC DNA].
MEDLINE=21435808; PubMed=11551509;
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                                                                                                                                                                Potently activated by enkephalins (By similarity).
SUBCELULIAR LOCATION: Integral membrane protein.
TISSUE SPECIFICITY: Uniquely localized in a subset of small dorsal root and trigeminal sensory neurons.
SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
                                                                                                                                                                                                                                                                                FUNCTION: Orphan receptor. Probably involved in the function nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain.
                                                                                                                                             Mas subfamily.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neurosci. 5:201-209(2002).
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1853733; PubMed=11850634; DOI=10.1038/nn815.
1.C., Grazzini E., Groblewski T., O'Donnell D., Roy M.-O.,
Hoffert C., Cao J., Schmidt R., Pelletier M., Labarre M.,
Hortin Y., Banville D., Shen S., Stroem P., Payza K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sensory neurons.";
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Zylka M.J., Simon M.I., Anderson D.J.;
                             no way modified and this statement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a new family of sensory
                                                     restrictions on
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MBL outstation -
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TRANSMEM
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                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      iGNC; HGNC:17617; MRGPRX4.
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                                                                                                                                                           PS00237; G_PROTEIN_RECEP_F1_1; 1.
PS50262; G_PROTEIN_RECEP_F1_2; 1.
n coupled receptor; Glycoprotein; Polymorphism; Receptor;
                             IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
                                                                                              CLSVLWPIWYRCRRPTHLSAVVCVLLWGLSLLFSMLEWRFCDFLFSGADSSWCETSDFIP
                                                                                                              CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT 180
                                                                                                                           VSIYILNLAAADFLFLSFQIIRSPLRLINISHLIRKILVSVMTFPYFTGLSMLSAISTER
VDKGEGQLPEESLELSGSRL
             VDEGGGWLPQETLELSGSKL
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                                                                                                                                                                                                                            322 AA;
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N-linked (GlcNAc...) (Pote
F -> L (in dbSNP:2468774).
/FTId=VAR_019435
N -> K (in dbSNP:2445180).
/FTId=VAR_019436
/FTId=VAR_019436
/FTId=VAR_019436
                                                                                                                                                                                                    Score 1366;
Pred. No. 6.
                                                                                                                                                                                                                                                           /FTId=VAR_019437.
S -> L (in dbSNP:2445179)
                                                                                                                                                                                                                                                                                                                                           Extracellular 7 (Potential)
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A -> V (in Ref. :
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RESULT

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Q5U9EO_MACFA
ID Q5U9EO, MACFA PRELIMINARY;
AC Q5U9EO;
AC MARMENIA;
AC CARROLLARIS (Crabe of Carcopithecidae; Cercopithecidae; Cercopithecox
AC ELMARYOTA; Metazoa; Chordat
AC MARMENIA; Eutheria; Euarcho
AC CECOPITACION AC ACTOPICA
AC ADAMA
AC ADA
RESULT 9
Q5U9D6 MACFA P
ID Q5U9D6 MACFA P
AC Q5U9D6,
DT 01-FEB-2005 (T
DT 01-FEB-200
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Best Local S
Matches 255
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PRINTS; PR00237; GFCREHODODSN.

PROSITE; PS00237; GPROTEIN RECEP F1 1; UNKNOWN 1.

PROSITE; PS50262; G_PROTEIN RECEP F1 2; 1.

G-protein coupled receptor; Receptor; Transducer; Transmembrane.

SEQUENCE 322 AA; 36781 MW; 54F5801F294BE07B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases.

-! SUBCELLULAR LOCATION: Integral membrane protein (By 18772457; AAV49124.1; -; Genomic DNA.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0001872; F:receptor activity; IEA.
GO; GO:0001874; F:rhodopsin-like receptor activity; IEA.
GO; GO:0007186; P:G-protein coupled receptor protein sign GO; GO:0007165; P:signal transduction; IEA.
InterPro; IPR00276; GPCR_Rhodpsn.
    Mas-related protein
Name=MrgX8;
Macaca fascicularis
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini;

Cercopithecidae; Cercopithecinae; Macaca.
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        (Crab
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79.2%;
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3 Receptors
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in Macaque, Mouse,
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Best Local S
Matches 244
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Q4QXU4;
13-SEP-2005 (
13-SEP-2005 (
13-SEP-2005 (
                                                                                                                                 Trachypithecus francoisi (Francois' langur) (Ine Bukaryota; Metazoa; Chordata; Craniata; Vertebra Mammalia; Butheria; Buarchontoglires; Primates; Cercopithecidae; Colobinae; Trachypithecus. NCBI_TaxID=54180;
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                                                                                                                                                                                                                                                                                           Name=MRGX2;
                                                                                                                                                                                                                                                                                                                         MRGX2.
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NG Receptors
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rimates; Catarrhini;
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in Macaque,
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NUCLEOTIDE SEQUENCE.
PubMed=15862286; DOI=10.1016/j.gene.2005.03.001;
Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L.,
"Adaptive evolution of MRGX2, a human sensory ne

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RESULT 11
Q4QXU2_PYGBI PRELIMINARY;
AC Q4QXU2;
AC Q4QXU2;
AC Q4QXU2;
DT 13-SEP-2005 (TrEMBLrel. 31,
DT MRGX2;
GN MCGATON MELAZOS; Chordate
OC Excaryota; Metazos; Chordate
OC Excaryota; Metazos; Chordate
OC Excaryota; Metazos; Chordate
OC Excaryota; Detheria; Exarchoto
OC Carcopithecidae; Colobinae;
OX NCBI_TAXID=61621;
RN NCCLEOTIDE SEQUENCE.
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Best Local S
Matches 210
  x PubMedd=15862286; DOI=10.1016/j.gene.2005.03.001;
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Involved in nociception.";

Gene 352C:30-35(2005).

C -!- SUBCELLULAR LOCATION: Integral membrane protein (By
EMBL; AY651166; AAM70079.1; -; Genomic_DNA.

InterPro; IFR000276; GPRC, Rhodpen.
R FAGN:TP: PF00001; Thm 1; 1.
R PF0001; Thm 1; 1.
R PROSITE; PR00237; GFCRHODOPSN.
R PROSITE; PS00237; GFROTEIN RECEP F1 1; UNKNOWN 1.
R PROSITE; PS00237; GFROTEIN RECEP F1 2; 1.
C SEQUENCE 330 AA; 37007 MW; F91875540B4716DE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Pygathrix bieti (Black snub-nosed monkey) (Rhinopithecus bieti). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Cercopithecidae; Colobinae; Pygathrix. NCBI_TaxID=61621;
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Pfam; PF00001; 7tm 1; 1.

PRINTS; PR00237; GPCRHODOPSN.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; UNKNOWN_1.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.

G-protein_coupled_receptor; Receptor; Transducer; TransCPCP_T2 = 1.

SEQUENCE 330 AA; 37043 MW; 4EE8ADD391CE443A CRC64;
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Gene 352C:30-35(2005).

-!- SUBCELLULAR LOCATION: Integral membrane
EMBL; AY651164; AAW70077.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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MRGX2_MACFA
Q5U9D9;
10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                    10-MAY-2005 (Rel. 47, Created)
10-MAY-2005 (Rel. 47, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
Mas-related G-protein coupled receptor member X2.
Name=MRGPRX2; Synonyms=MRGX2;
Macaca fascicularis (Crab eating macague) (Cynomolgus monkey).
                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
Zhang L., Taylor N., For "Cloning and expression"
                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebro
Mammalia; Eutheria; Euarchontoglires; Primates;
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NCBI_TaxID=9541;
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                                                                                                                                                    s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no restr as long as its content is in no way modified and this s
                                                                                                                                                                                                                     SUBCELLULAR LOCATION: Integral membrane protein. SIMILARITY: Belongs to the G-protein coupled rec
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Cytoplasmic (Potential).
2 (Potential).
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Matches 206
-!- SUBCELLULAR LOCATION: Integral membrane premate, AY651163, AAW70076.1; -; Genomic_DNA.
Interpro; IPR000276; GPCR Rhodpsn.
Pfam; PF00001; 7tm 1; 1.
PRINTS; PR00237; GPCRHODOPSN.
PROSITE; PS00237; GPCRHODOPSN.
PROSITE; PS00237; GPROTEIN RECEP_F1_1; UNKNOW PROSITE; PS50262; G_PROTEIN RECEP_F1_2; 1.
G-protein coupled receptor; Receptor; Transductive of the coupled receptor of the coupled
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Q4QXU5;
13-SEP-2005 (
13-SEP-2005 (
13-SEP-2005 (
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Macaca mulatta (Rhesus macaque).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini;
Cercopithecidae; Cercopithecinae; Macaca.
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Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao
"Adaptive evolution of MRGX2, a human sensory neuron
                                                                                                                                                                                                                                                                                                                                                                                                                                              Cercopithecidae;
NCBI_TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=MRGX2;
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ilarity 62.6%;
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Pred. No. 2.5e
29; Mismatches
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Cytoplasmic (Potent; 6C783BDBC7EC246E
     02AB400A7BC7EB99 CRC64;
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                             Transducer; Transmembrane
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Best Local Similarity
Matches 209; Conser
                                                                                                                                                                                                       Submitted (JIT--
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25-OCT-2004
25-OCT-2004
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausher R.D., Collins F.S., Wagner U.S., Shenmen C.M., Schuler (Klausher R.D., Collins F.S.) Betow K.H., Schaefer C.F., Bhat N.I Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.I Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
                                                                                                                                                                                                                                                                       Suwa M., Sato T., Okouo
Tsutsumi S., Aburatani
"Genome-wide discovery
                                                                                                                                                                                                                                                                                                                                                                                                   genome sequence.";
FEBS Lett. 520:97-101(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=22040266; Pub
Takeda S., Kadowaki S
"Identification of G
genome sequence.";
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25-OCT-2004 (Rel. 45, Last sequence update)
25-OCT-2005 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
Mas-related G-protein coupled receptor member X2.
Name=MRGPRX2; Synonyms=MRGX2;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nociceptive sensory neurons."; Cell 106:619-632(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=21435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4; Dong X., Han S.-K., Zylka M.J., Simon M.I., Anderson D.J.; "A diverse family of GPCRs expressed in specific subsets of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MRGX2
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                                                                                                                                                                                                                                                                  SEQUENCE.
iato T., Okouchi I., Arita M., Futa
3., Aburatani H., Asai K., Akiyama
3., Aburatani H., and analysis of human
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435808; PubMed=11551509; DO1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PubMed=12044878; DOI=10.1016/S0014-5793(02)02775-8;
ki S., Haga T., Takaesu H., Mitaku S.;
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EMBL; AY042214; AAK91805.1; -; Genom EMBL; AB083626; BAB89339.1; -; Genom EMBL; AB065811; BAC06030.1; -; Genom EMBL; BC063450; AAH63450.1; -; mRNA. Ensembl; ENSG00000183695; Homo sapie HGNC; HGNC:17983; MRGPRX2.

MIM; 607228; -. C:integral to membra GO; GO:0016021; C:integral to membra GO; GO:0004930; F:G-protein coupled GO; GO:0004931; P:neuropeptide bindi GO; GO:001923; P:perception of pain GO; GO:0030431; P:sleep; NAS.

InterPro; IPR000276; GFCR_Rhodpsn.
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Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length hum.
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J. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=12915402; DOI=10.1074/jbc.M302456200; Robas N., Mead E., Fidock M.;
                                                                                                                                                                  Pfam; PFUUUU1; '.'" C'RHODOPSN.
PRINTS; PR00237; GĒCRHODOPSN.
PROSITE; PS00237; G PROTEIN RECEP F1 1; 1.
PROSITE; PS50262; G PROTEIN RECEP F1 2; 1.
PROSITE; PS50262; G PROTEIN RECEP F1 3; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
                                                                                                                                                                                                                                                        Pfam; PF00001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           removed
                                                                                                                                                                                                                                                                                     GO:0016021; C:integral to membrane; IC. GO:0016021; C:integral to membrane; IC. GO:0004930; F:G-protein coupled receptor GO:0042933; F:neuropeptide binding; IPI. GO:0019233; P:perception of pain; NAS. GO:0030431; P:sleep; NAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coupled.
SUBCELLULAR LOCATION: Integral membrane protein.
SUBCELLULAR LOCATION: Integral membrane protein.
TISSUE SPECIFICITY: Has a limited expression profile, both
perispheral and within the central nervous system, with highest
levels in dorsal root ganglion.
SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FUNCTION: Orphan receptor. Probably involved in the function nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Cortistatin-14 seems to be a high potency ligand at this reconstitution has several biological functions including roles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           European Bioinformatics Institute. as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mas subfamily.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sleep regulation locomotor activity, and cortical function. In receptor-expressing cells, cortistatin-stimulated increases in intracellular Ca(2+) but had no effect on basal or forskolinstimulated cAMP levels, suggesting that this receptor is G(q)-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lion."
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Toshiyuki S., Carninci P.,
Peters G.J., Abramson R.D.,
         1 (Potential).

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2 (Potential).
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Mullahy S.J.,
naratne P.H.,
J., Hulyk S.W.,
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MBL outstation -
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Best Local &
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  involved in nociception.";

Gene 352:30-35(2005).

-i- SUBCELLULAR LOCATION: Integral membrane pr
-i- SUBCELLULAR LOCAT
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13-SEP-2005 (TrEMBLrel. 3:
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Eukaryota; Metazoa; (
Mammalia; Eutheria; E
                                                                                                                                                                                                                                                   Yang S., Liu Y., Lin A.A., Ca
"Adaptive evolution of MRGX2,
                                                                                                                                                                                                                                                                                                                                                                                                                        Yang S., Liu Y., Lin A.A., Cav
"Adaptive evolution of MRGX2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PubMed=15862286; DOI=10.1016/j.gene.2005.03.001; Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE
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HUMAN
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K2, a human sensory neuron
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APPLICANT: Peri, Krishna G.

APPLICANT: Moffett, Serge

APPLICANT: Moffett, Serge

APPLICANT: Moffett, Serge

APPLICANT: Moffett, Serge

APPLICANT: Moffett, Serge

APPLICANT: MOFFERD AND COMPOUNDS FOR PREVENTION AND TREATMENT OF ELEVATED

TITLE OF INVENTION: INTRACCULAR PRESSURE AND RELATED CONDITIONS

FILE REFERENCE: 4518/1M674US1

FULE REFERENCE: 4518/1M674US1

FULE REFERENCE: 4518/1M674US1

CURRENT FILING DATE: 2003-03-27

CURRENT FILING DATE: 2002-03-27

PRIOR APPLICATION NUMBER: US 60/367,513

PRIOR APPLICATION NUMBER: US 60/367,513

PRIOR FILING DATE: 2002-03-27

NUMBER OF SEQ ID NOS: 8

SOFTWARE: PatentIn version 3.1

SEQ ID NO 2

LENGTH: 322

TURNER: DETERMINED AND TREATMENT OF ELEVATED.
SOFTWARE: E
SEQ ID NO 5
LENGTH: 322
                                                                         GENERAL INFORMATION:

APPLICANT: Ahmad, Sultan
APPLICANT: Banville, Denis
APPLICANT: Banville, Denis
APPLICANT: Comment of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the service of the
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ORGANISM: Homo sapiens
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Pred. No. 3.7e-133;
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SOFTWARE: Patentin vers
SEQ ID NO 7
LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
US-09-254-227A-7
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US-09-254-227A-7
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Best Local Similarity 94.1%;
Matches 302; Conservative
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Best Local (
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APPLICANT: Lembo, Paola
APPLICANT: Lembo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from
FILE REFERENCE: 9.8823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
NUMBER OF SEQ ID NOS: 22
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APPLICANT: Banville,
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                                  IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
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82.3%; Pred. No. 2.5e-112;
tive 24; Mismatches 33;
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Pred. No. 1.8e-127;
8; Mismatches 11;
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Sequence 9, Application US/09254227A

Patent No. 6696257

GENERAL INFORMATION:

APPLICANT: Ahmad, Sultan
APPLICANT: Banville, Denis
APPLICANT: Hembo, Paola
APPLICANT: Combo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Haiang, Shen
ITILE OF INVENTION: G Protein-Coupled Receptors from the
FILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn version 3.0
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US-10-314-048A-20
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SEQ ID NO 9
LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
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Sequence 20, Application US/10314048A
Patent No. 6902902
GENERAL INFORMATION:
APPLICANT: Unett, David J.
APPLICANT: Chen, Ruoping
APPLICANT: Richman, Jeremy
APPLICANT: Connolly, Daniel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      81.7%; Score 1381; DB 2
82.3%; Pred. No. 4e-111;
cive 22; Mismatches 3
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APPLICANT: Lowitz, Kevin P.
APPLICANT: Behan, Dominic P.
APPLICANT: Behan, Dominic P.
APPLICANT: Chalmers, Derek T.
APPLICANT: Lerner, Michael
TITLE OF INVENTION: Human G Protein-Coupled Receptors and Modulators Thereof
TITLE OF INVENTION: for the Treatment of Metabolic-Related Disorders
FILE REFERENCE: 22.USS.CIP
CURRENT APPLICATION NUMBER: US/10/314,048A
CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: 10/996,511
PRIOR APPLICATION NUMBER: 00/995,543
PRIOR APPLICATION NUMBER: 00/399,917
PRIOR APPLICATION NUMBER: 60/399,917
PRIOR APPLICATION NUMBER: 60/399,917
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR FILING DATE: 2002-09-13
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR FILING DATE: 2002-09-13
NUMBER: 07-507 ID NOSE: 50/410,747
PRIOR FILING DATE: 2002-09-13
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US-09-254-227A-13
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 20
LENGTH: 322
Type: no.
Sequence 13, Application US/09254227A

Patent No. 6696257

GENERAL INFORMATION:
APPLICANT: Ahmad, Sultan
APPLICANT: Banville, Denis
APPLICANT: Fortin, Yves
APPLICANT: Lembo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 81.2%; Score 1373; DB 2; Best Local Similarity 82.0%; Pred. No. 1.9e-110; Matches 264; Conservative 22; Mismatches 36;
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Liaw, Chen
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rd, James
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Rat and

Human

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; SOFTWARE: PatentIn veri
; SEQ ID NO 11
; LENGTH: 322
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-254-227A-11
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US-09-254-227A-11
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SEQ ID NO 13
LENGTH: 322
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Best Local Similarity
Matches 264; Conserv
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CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
CORTUNNES 22
                                                                                                                                                                                                                                                                              FILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                                                                                                                                                              APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from the Rat and Human
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                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ahmad, Sultan APPLICANT: Banville, Den
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Local Similarity 83.1%;
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MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
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Lembo, Paola
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                                                                                                                ; Score 1356; DB 2;
; Pred. No. 5.6e-109;
21; Mismatches 35;
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Pred. No. 9.4e-110;
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PRIOR FILING DATE: 2002-05-13
PRIOR APPLICATION NUMBER: 60/410,747
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 161
SOFTWARE: PatentIn version 3.1
SEQ ID NO 30
LENGTH: 330
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US-10-314-048A-30
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                                                                                                                                                                                Query Match 57.9%; Score 978.5; DB 2; Best Local Similarity 62.3%; Pred. No. 1.7e-76; Matches 205; Conservative 29; Mismatches 86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/314,048A CURRENT FILING DATE: 2002-12-06 FRIOR APPLICATION NUMBER: 10/096,511 PRIOR FILING DATE: 2002-03-12 PRIOR APPLICATION NUMBER: 09/995,543 PRIOR FILING DATE: 2001-11-27 PRIOR FILING DATE: 2001-11-27 PRIOR FILING DATE: 2001-11-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Lowitz, Kevin P.
APPLICANT: Behan, Dominic P.
APPLICANT: Chalmers, Derek T.
APPLICANT: Chalmers, Michael
TITLE OF INVENTION: Human G Protein-Coupled Receptors and Modulators Thereof
TITLE OF INVENTION: for the Treatment of Metabolic-Related Disorders
FILE REFERENCE: 22.US6.CIP
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APPLICANT:
APPLICANT:
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PRIOR FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR FILING DATE: 2002-08-19
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                                    RNAFSVYVLSLAGADFLFLCFQIINCLVYLSNFFCSISINFPSFFTTVMTCAYLAGLSML
                                                          RNAVSIYILNLVAANFLFLSGHIIFSPLFLINIRHPIS----KILSPVMTFPYFIGLSML
                                                                                                              MDPTTPAWGTESTTVNGNDQALLLLCGKETLIPVFLILFIALVGLVGNGFVLWLLGFRMR
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Leonard, James
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Liaw, Chen
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APPLICANT: Ahmad, Sultan
APPLICANT: Banville, Denis
APPLICANT: Fortin, Yves
APPLICANT: Lembo, Paola
APPLICANT: Lembo, Paola
APPLICANT: Lembo, Paola
APPLICANT: Shi-Hsiang, Shen
APPLICANT: Shi-Hsiang, Shen
CURRENT SHI-HSIANG, Shen
FILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1
LENGTH: 337
TYPE: PRT
ORGANISM: rat
ORGANISM: rat
              US-10-314-048A-10

US-10-314-048A-10

; Sequence 10, Application US/10314048A

; Patent No. 6902902
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US-09-254-227A-1
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Patent No. 6696257
GENERAL INFORMATION:
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 173;
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                                                                                                                                         RALODTPEVDEGGGWLPQETLELS 316
                                                                                                                                                                                           QWAL--FSRIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQ
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             OLIAVIO; OTTALE OF INVENTION: DNA SEQUENCES ENCODING MAS ONHCOGENE; POLYPETIDES ENCODED THEREFROM AND DIAGNOSTIC AND OTHER METHODS
BASED THEREFROM
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APPLICANT:
APPLICANT:
                                                           APPLICANT: Young, Dallan; Wigler, Michael H.; Fasano
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CURRENT APPLICATION NUMBER: US/10/314,048A CURRENT FILING DATE: 2002-12-06

PRIOR APPLICATION NUMBER: 10/096,511

PRIOR FILING DATE: 2002-03-12

PRIOR PAPPLICATION NUMBER: 09/995,543

PRIOR FILING DATE: 2001-11-27

PRIOR APPLICATION NUMBER: 60/399,917

PRIOR FILING DATE: 2002-07-29

PRIOR APPLICATION NUMBER: 60/404,761

PRIOR APPLICATION NUMBER: 60/404,761

PRIOR APPLICATION NUMBER: 60/404,761

PRIOR APPLICATION NUMBER: 60/410,747

PRIOR APPLICATION NUMBER: 60/410,747

PRIOR FILING DATE: 2002-09-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 30.2%; Score 510; DB 2; Length 321; Best Local Similarity 40.3%; Pred. No. 3.4e-36; Matches 127; Conservative 57; Mismatches 115; Indels
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SOFTWARE: Patentin version 3.1
SEQ ID NO 10
LENGTH: 321
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APPLICANT: Chen, R
APPLICANT: Richman
APPLICANT: Connoll
APPLICANT: Dang, H
APPLICANT: Choi, B
APPLICANT: Leonard
APPLICANT: Hakak,
APPLICANT: Liaw, C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Chalmers, Derek T.
APPLICANT: Lerner, Michael
TITLE OF INVENTION: Human G Protein-Coupled Receptors and Modulators Thereof
TITLE OF INVENTION: for the Treatment of Metabolic-Related Disorders
FILE REFERENCE: 22.US6.CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
                                                                                         180 MVQAALIMGVLTEVMTLSSLTLEVMVRRSSQQWRRQP-TRLEVVVLASVLVFLICSLPLS
                                                                                                                                                                                                                                                                                                              118 TERCLSILMPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVMCETSD
                                            291 LORALODTPEVDEGG 305
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LOOALREEPEL-EGG
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Choi, Bryan
Leonard, James
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Liaw, Chen
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Connolly, Daniel
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NUMBER OF SEQUENCES:

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RESULT 13
US-08-118-270-52
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APPLICATION NUMBER: US/
FILING DATE: 06-JUN-1986;
SEQ ID NO:2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 24.6
Best Local Similarity 34.1
Matches 105; Conservative
                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/118,270

FILING DATE: 09-SEP-1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/943,236

FILING DATE: 10-SEP-1992

ATTORNEY/AGENT INFORMATION:

NAME: TOWNSEND, NUMBER: 34,033

REFERENCE/DOCKET NUMBER: MURPHY=2A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 5508384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Murphy, Randall B. APPLICANT: Schuster, David I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
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TELEPHONE: 202-737-3528
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ZIP: 20004
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                                                                                                                                                                                                                                                                                                                                                                                                                                        E: BROWDY AND NEIMARK
419 Seventh Street, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 348
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Best Local |
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                                                                                                                                                        PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/943,236

FILING DATE: 10-SEP-1992

ATTORNEY/AGENT INFORMATION:

NAME: Townsend, Kevin G.

REGISTRATION NUMBER: 34,033

REFERENCE/DOCKET NUMBER: MURPHY=2 P.
                                                                    TELEX: 248633
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: New York University
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
KUMBER OF SEQUENCES: 348
CORRESPONDENCE ADDRESS:
ADDRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
COURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08528
                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
                             SEQUENCE CHARACTERISTICS LENGTH: 282 amino aci
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 22.9%; Score 386.5; DB 1; Local Similarity 34.4%; Pred. No. 1.3e-25;
                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 20004
                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
STRANDEDNESS:
                                                                                                                                                                                                                                                                                     FILING DATE: 09-SEP-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     282 amino acids
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419 Seventh Street, N.W.,
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                                                                                                                                                              MURPHY=2 PCT
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; TOPOLOGY: linear ; MOLECULE TYPE: peptide US-08-118-270-76
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US-08-118-270-76
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Best Local Similarity 34.4
Matches 96; Conservative
                                                                                        TELEX: 248633
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 298 amino acid
                                                                                                                                                                                                                                                                                      SOFTWARE: PACENTIN Release #1.0, Version CURRENT APPLICATION NUMBER: US/08/118,270 FILLING DATE: 09-SEP-1993 PRIOR APPLICATION NUMBER: US 07/943,236 PILLING DATE: 10-SEP-1992 PILLING DATE: 10-SEP-1992 PATORNEY/AGENT INFORWATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Murphy, Randall B.

APPLICANT: Schuster, David I.

TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN

TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND M
                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
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                                                                                                                                                           TELEPHONE: 202-0-3528
                                                                                                                                                                                                                                   NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY=2A
                                                     TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                            298 amino acids
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419 Seventh Street, N.W.,
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      125 LLSFLVTSIHNYFC-LLGHEASGTACLNMDISLLGILLFFLFCPIMVLPC----IALLHV 179
                                                                                                                                                                                                                                            234 CICINSSAKPIVYFIAGRDKSORLWEPLRVVFORALRDGAEPGDAASSTPNTVTMEM
                                                                                                                                                                                                                                                                                                                                                                                                                   260 LSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQE-TLEL 315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       203 LCGSRKMPLT-RLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDW--KVLFCHVHLVSIF 259
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RESULT US-10-2 US-10-2 Seque Publi GENER GENER APPL APPL APPL APPL FILE FILE FILE FILE FILE FILE FILE FIL	RESULT 1 US-10-292-798-1274 Sequence 1274, Application US/1029 PUBLICATION NO. US20030235833A1 GENERAL INFORMATION: APPLICANT: SUWA, MAKIKO APPLICANT: ASAL, KIYOSHI APPLICANT: ASAL, KIYOSHI APPLICANT: ASURATANI, HIROYUKI INTILE OF INVENTION: GUANOSINE TRI FILE REFERENCE: 084335/166 CURRENT APPLICATION NUMBER: US/10 FILE REPERENCE: 084335/166 CURRENT FILING DATE: 2002-11-13 PRIOR FILING DATE: 2001-12-18 PRIOR FILING DATE: 2001-12-18 PRIOR FILING DATE: 2001-06-18 NUMBER OF SEQ ID NOS: 2070 SOFTWARE: PatentIN Ver. 2.1 SEQ ID NO 1274 LENGTH: 322	ALIGNMENTS 92798 161 1-246789	PROTEIN COUPLED RECE
Query Me Best Loc Matches Qy	atch bal S 314	97.4%; Score 1647; DB 4; Length 322; iimilarity 97.5%; Pred. No. 8.9e-140; Conservative 4; Mismatches 4; Indels 0; Gaps MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA	Length 322; ; Indels 0; Gaps 0; LVALTGNAVVLWLLGCRMRRNA 60
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APPLICANT: Rieger, Daniel K.
APPLICANT: Burgess, Catherine E.
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Sam
FILE REFERENCE: 21402-258
CURRENT APPLICATION NUMBER: US/10/072,012
CURRENT FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: 60/265,102
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-31
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NUMBER OF SEQ ID NOS: 1391
SOSTWARE: PatentIn Ver. 2.1
SEQ ID NO 530
LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
US-10-072-012-530
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                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 2001-02-08
                                                                                          314;
MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
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Patturajan, Muu-
Patturajan, Muu-
Patta, Richard
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Rastelli,
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Rieger, Daniel K.
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Grosse, William M.
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Gusev, Vladimir Y.
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Zerhusen, Bryan
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5. US20040033493A1
                                                                                          97.4%;
llarity 97.5%;
Conservative
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David W.
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                                                                                          Score 1647; DB 4;
Pred. No. 8.9e-140;
4; Mismatches 4;
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CURRENT APPLICATION NUMBER: US/10/072,012
CURRENT FILING DATE: 2002-01-31
CURRENT FILING DATE: 60/265,102
PRIOR APPLICATION NUMBER: 60/265,514
PRIOR APPLICATION NUMBER: 60/265,514
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PRIOR APPLICATION NUMBER: 60/265,517
PRIOR FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 60/265,412
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APPLICANT: Rieger, Daniel K.
APPLICANT: Burgess, Catherine E.
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding
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DR FILING DATE: 2001-02-02

PR APPLICATION NUMBER: 60/266,767

PR FILING DATE: 2001-02-05

PR APPLICATION NUMBER: 60/267,057

PR FILING DATE: 2001-02-07

PR APPLICATION NUMBER: 60/266,975

PR FILING DATE: 2001-02-07

PR APPLICATION NUMBER: 60/267,459

PR APPLICATION NUMBER: 60/267,459
                                                                                                                                      FILING DATE: 2001-01-31
APPLICATION NUMBER: 60/265,395
FILING DATE: 2001-01-31
APPLICATION NUMBER: 60/266,406
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Gerlach, Valerie
Taupier Jr, Raymond J
Gusev, Vladimir Y.
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Grosse, William M.
Alsobrook II, John P.
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Patturajan, Mee
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TITLE OF INVENTION: Endogenous And No. US20020193584A1-Endogenous Versititle OF INVENTION: Receptors
FILE REFERENCE: AREN-0308
CURRENT APPLICATION NUMBER: US/09/995,225
CURRENT FILING DATE: 2001-11-26
PRIOR APPLICATION NUMBER: 09/170,496
PRIOR FILING DATE: 1998-10-13
PRIOR APPLICATION NUMBER: PCT/US99/23938
PRIOR APPLICATION NUMBER: PCT/US99/23938
PRIOR APPLICATION NUMBER: PCT/US99/23938
PRIOR APPLICATION NUMBER: 60/253,404
PRIOR FILING DATE: 1998-10-13
PRIOR APPLICATION NUMBER: 60/253,404
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/255,366
PRIOR APPLICATION NUMBER: 60/270,286
PRIOR APPLICATION NUMBER: 60/270,286
PRIOR APPLICATION NUMBER: 60/270,286
PRIOR APPLICATION NUMBER: 60/282,365
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NUMBER OF SEQ ID NOS: 1391
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 535
LENGTH: 322
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            OR FILING DATE: 2001-04-06
OR APPLICATION NUMBER: 60/270,266
OR FILING DATE: 2001-02-20
OR APPLICATION NUMBER: 60/282,032
OR FILING DATE: 2001-04-06
OR APPLICATION NUMBER: 60/282,358
OR FILING DATE: 2001-04-06
OR APPLICATION NUMBER: 60/282,356
OR APPLICATION NUMBER: 60/282,356
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Local Similarity 97.5%;
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Chu, Zhi Liang
Dang, Huong T.
Lowitz, Kevin P.
Pride, Cameron
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Pred. No. 8.9e-140;
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PILE REFERENCE: AREN-0308

CURRENT APPLICATION NUMBER: US/09/995,225

CURRENT FILING DATE: 2001-11-26

PRIOR APPLICATION NUMBER: 09/170,496

PRIOR FILING DATE: 1998-10-13

PRIOR APPLICATION NUMBER: PCT/US99/23938

PRIOR FILING DATE: 1998-10-13

PRIOR FILING DATE: 1998-10-13

PRIOR FILING DATE: 1998-10-13

PRIOR APPLICATION NUMBER: 60/253,404

PRIOR APPLICATION NUMBER: 60/253,366

PRIOR APPLICATION NUMBER: 60/255,366
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PRIOR FILING DATE: 2001-05-14
PRIOR APPLICATION NUMBER: 60/309,208
PRIOR FILING DATE: 2001-07-31
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.1
SEQ ID NO 20
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Best Local :
                                                                                                                                                                                                                                                                                                                         APPLICANT: LOWİtz, Kevin P. APPLICANT: Pride, Cameron TITLE OF INVENTION: Endogenous TITLE OF INVENTION: Receptors
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ORGANISM: Artificial Sequence
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DR FILLING DATE: 2000-12-12

DR APPLICATION NUMBER: 60/270,286

DR FILLING DATE: 2001-02-20

DR APPLICATION NUMBER: 60/282,365

DR FILLING DATE: 2001-04-06

DR APPLICATION NUMBER: 60/270,266

DR APPLICATION NUMBER: 60/270,266

DR APPLICATION NUMBER: 60/270,266

DR APPLICATION NUMBER: 60/270,266

DR APPLICATION NUMBER: 60/270,266
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                                                                                                                                                                                                                                                                                                                                                                                                    Dang, Huong T.
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Chu, Zhi Liang
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Pred. No. 2.5e-139;
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Sequence 31, Application US/10183116

Publication No. US20030092035A1

GENERAL INFORMATION:
APPLICANT: Anderson, David J.
APPLICANT: Dong, Xinzhong
APPLICANT: Sylka, Mark
APPLICANT: Sylka, Mark
APPLICANT: Sylka, Mark
APPLICANT: Han, Sang-kyou
TITLE OF INVENTION: PAIN SIGNALING MOLECULES
TITLE OF INVENTION: PAIN SIGNALING MOLECULES
TITLE OF APPLICATION NUMBER: US/10/183,116
CURRENT APPLICATION NUMBER: US 60/222,344
PRIOR APPLICATION NUMBER: US 60/222,344
PRIOR FILING DATE: 2000-08-01
PRIOR APPLICATION NUMBER: US 09/20,027
PRIOR APPLICATION NUMBER: US 09/704,707
PRIOR FILING DATE: 2000-11-03
PRIOR FILING DATE: 2000-10-05-04
PRIOR PRIOR APPLICATION NUMBER: US 09/704,707
PRIOR FILING DATE: 2001-04-19
PRIOR APPLICATION NUMBER: US 09/849,869
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
PRIOR PRILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
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PRIOR APPLICATION NUMBER: 60/282,358
PRIOR APPLICATION NUMBER: 60/282,356
PRIOR APPLICATION NUMBER: 60/282,356
PRIOR APPLICATION NUMBER: 60/290,917
PRIOR PILING DATE: 2001-05-14
PRIOR PILING DATE: 2001-05-14
PRIOR PILING DATE: 2001-07-31
NUMBER: OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.1
SEQ ID NO 20
LENGTH: 322
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Best Local Similarity 97.4
313; Conservative
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97.2%;
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Pred. No. 2.5e-139;
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; ORGANISM: Homo
US-10-183-116-31
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Best Local S
Matches 313
                                                                                                                                                                                                                                                                                        APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 1920-4-4
CURRENT FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/57,144
PRIOR APPLICATION NUMBER: 60/57,144
PRIOR FILING DATE: 2000-12-19
NUMBER: 0F SEQ ID NOS: 2292
SOFTWARE: PATENTIN Version 3.1
SEQ ID NO 674
LENGTH: 322
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SEQ ID NO 31
LENGTH: 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 674, Applic Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                Matches 313;
                                                                                                                                                                                               Query Match
Best Local Similarity
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121 CLSILMPIMYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
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                                                                 61 VSIYILNIVAANFLFLSGHIIFSPLFLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
                                                                                                                                   1 MDPTIPVLGTKLTPINGREBTPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRWRRNA
                                                                                                                  1 MDSTIPVLGTELTPINGREETPCYKQTLSFTGLTCIVSLVALTGNAVVLWLLGCRMRRNA
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                                                  VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
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o. US20030113798A1
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97.2%;
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                                                                                                                                                                                             Score 1642; DB 4;
Pred. No. 2.5e-139;
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Pred. No. 2.5e-139;
                                                                                                                                                                                  Mismatches
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RESULT 8 US-10-072-012-529

Sequence 529, Appropriate Publication No.

GENERAL INFORMATION: APPLICANT:

Tchernev,

Velizar

Patturajan, Zerhusen, Bryan Spytek, Kimberly Ś ₽ δ

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181

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APPLICANT: Burgess, Catherine E.
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Sam
FILE REFERENCE: 21402-258
CURRENT APPLICATION NUMBER: US/10/072,012
CURRENT APPLICATION NUMBER: 60/265,102
PRIOR APPLICATION NUMBER: 60/265,102
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/265,514
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR FILING DATE: 2001-01-31
PRIOR FILING DATE: 2001-01-31
PRIOR PRILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR PRILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR APPLICATION NUMBER: 60/266,406
PRIOR FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR PILING DATE: 2001-02-07
PRIOR PILING DATE: 2001-02-07
PRIOR PILING DATE: 2001-02-07
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PRIOR PILING DATE: 2001-02-08
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PRIOR PILING DATE: 2001-02-08
                                                                        Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 1391
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Grosse, William M.
Alsobrook II, John P.
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Rastelli, Luca
Miller, Charles E.
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Rieger, Daniel K.
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b. US20040033493A1
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/ladimir Y.
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            APPLICANT: Burgess, Catherine E.
TITLE OF INVENTION: Proteins and Nucleic Acids Engriller Reperrence: 21402-258
CURRENT APPLICATION NUMBER: US/10/072,012
CURRENT FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: 60/265,102
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/265,514
PRIOR APPLICATION NUMBER: 60/265,514
PRIOR FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR FILING DATE: 2001-01-31
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PRIOR FILING DATE: 2001-01-31
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US-10-072-012-534
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Matches 313
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FILING DATE:
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Taupier Jr, Raymond J
Gusev, Vladimir Y.
Colman, Steven D.
Wolenc, Adam R.
Pena, Carol B. A
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Spytek, Kimberly
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Shimkets, Richard
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Rieger, Daniel K.
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Grosse, William M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rastelli,
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Anderson, David W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gangolli, Esha
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97.2%;
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Pred. No. 2.5e-139;
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APPLICANT: APPLICANT:
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Gerlach, Valerie

Gusev, Vladimir Taupier APPLICANT:

Padigaru,

Esha

SOFTWARE: Pat SEQ ID NO 529

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PRIOR APPLICATION NUMBER: 60/266,406
PRIOR FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR APPLICATION NUMBER: 60/267,057
PRIOR APPLICATION NUMBER: 60/267,057
PRIOR APPLICATION NUMBER: 60/267,057
PRIOR APPLICATION NUMBER: 60/267,975
PRIOR APPLICATION NUMBER: 60/266,975
PRIOR FILING DATE: 2001-02-07
PRIOR FILING DATE: 2001-02-07
PRIOR FILING DATE: 2001-02-08
Remaining Prior Application data removed - See File Wrapper or PALA
NUMBER OF SEQ ID NOS: 1391
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 534
LENGTH: 322
APPLICANT: 2ylka, Mark
APPLICANT: Simon, Melvin
APPLICANT: Simon, Melvin
APPLICANT: Han, Sang-kyou
FIITLE OF INVENTION: PAIN SIGNALING MOLECULES
FILE REFERENCE: CALTE.4C1CP1C1
CURRENT APPLICATION NUMBER: US/10/957,135
CURRENT FILING DATE: 2004-09-30
PRIOR APPLICATION NUMBER: US 60/222,344
PRIOR APPLICATION NUMBER: US 60/202,027
PRIOR FILING DATE: 2000-08-01
PRIOR PILING DATE: 2000-05-04
PRIOR PILING DATE: 2000-11-03
PRIOR PILING DATE: 2000-11-03
PRIOR APPLICATION NUMBER: US 60/285,493
PRIOR APPLICATION NUMBER: US 60/285,493
PRIOR FILING DATE: 2001-04-19
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
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US-10-957-135-31
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; ORGANISM: Homo sapiens
US-10-072-012-534
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Best Local Similarity 9/...
13; Conservative
                                                                                                                                                                                                                                                                                                                                                               Sequence 31, Application US/10957135
Publication No. US20050037468A1
GENERAL INFORMATION:
APPLICANT: Anderson, David J.
APPLICANT: Dong, Xinzhong
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97.2%;
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Pred. No. 2.5e-139;
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; PRIOR APPLICATION NUMBER: US 10/183,116
; PRIOR FILING DATE: 2002-06-26
; NUMBER OF SEQ ID NOS: 109
; SOPTWARE: FASTSEQ for Windows Version 4.
; SEQ ID NO 31
; SEQ ID NO 31
; LENGTH: 322
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-957-135-31
                                                                                                                                                   FILE REFERENCE: CALTE .004C1

CURRENT APPLICATION NUMBER: US/11/083,611

CURRENT FILING DATE: 2005-03-17

PRIOR APPLICATION NUMBER: US 09/849,869

PRIOR FILING DATE: 2001-05-04

PRIOR APPLICATION NUMBER: US 60/222,344

PRIOR FILING DATE: 2000-08-01

PRIOR APPLICATION NUMBER: US 60/202,027

PRIOR FILING DATE: 2000-05-04

PRIOR FILING DATE: 2000-11-03

PRIOR APPLICATION NUMBER: US 60/202,027

PRIOR APPLICATION NUMBER: US 60/202,027

PRIOR APPLICATION NUMBER: US 60/203,493

PRIOR FILING DATE: 2000-11-03

PRIOR PILING DATE: 2001-04-19

NUMBER OF SEQ ID NOS: 115
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                                                     US-11-083-611-31
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Best Local S
Matches 313
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                                                                                                   SOFTWARE: FastSEQ for Windows Version SEQ ID NO 31 LENGTH: 322
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Anderson, David J.
APPLICANT: Dong, Xinzhong
APPLICANT: Zylka, Mark
APPLICANT: Simon, Melvin
APPLICANT: Simon, Melvin
APPLICANT: Han, Sang-Kyou
TITLE OF INVENTION: PAIN SIGNALING MOLECULES
                                                                     TYPE: PRT
ORGANISM: Homo sapiens
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Pred. No. 2.5e-139;
5; Mismatches 4;
Score 1642; DB 6;
Pred. No. 2.5e-139;
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Query Match Best Local Similarity

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322

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: WEI, Ming-Hui et al.
APPLICANT: WEI, Ming-Hui et al.
TITLE OF INVENTION: ISOLATED HUMAN G-PROTEIN COUPLED
TITLE OF INVENTION: RECEPTORS, NUCLEIC ACID MOLECULES ENCY
TITLE OF INVENTION: RECEPTORS, AND USES THEREOF
FILE REFERENCE: CL000900-CIP
CURRENT APPLICATION NUMBER: US/09/867,570
CURRENT APPLICATION NUMBER: 09/695,045
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 09/695,045
PRIOR FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 337
TYPE: PRT
ORGANISM: Human
US-09-867-570-2
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                    RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                               IAMLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
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 RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                               CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGADSVWCETSDFIT
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Sequence 2, Application US/10401397A

Publication No. US20030212001A1

GENERAL INFORMATION:

APPLICANT: Peri, Krishna G.

APPLICANT: Moffett, Serge

APPLICANT: Moffett, Serge

ITILE OF INVENTION: METHODS AND COMPOUNDS FOR PRE

TITLE OF INVENTION: INTRAOCULAR PRESSURE AND REI

FILE REFERENCE: 4518/1M674US1

CURRENT APPLICATION NUMBER: US/10/401,397A

CURRENT FILING DATE: 2003-03-27

PRIOR APPLICATION NUMBER: US 60/367,513

PRIOR FILING DATE: 2002-03-27

NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Takeda Chemical Industries, Ltd.
TITLE OF INVENTION: Determination of a ligand
FILE REFERENCE: P03-0006PCT
CURRENT APPLICATION NUMBER: US/10/505,486
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: UP 2002-45728
PRIOR FILING DATE: 2002-02-22
PRIOR FILING DATE: 2002-02-22
PRIOR PFLICATION NUMBER: UP 2002-213949
PRIOR FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: UP 2002-298237
PRIOR APPLICATION NUMBER: UP 2002-298237
PRIOR FILING DATE: 2002-10-11
NUMBER OF SEQ ID NOS: 233
SEQ ID NO 104
LENCTH: 560
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Best Local S
Matches 313
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Pred. No. 4.5e-139;
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Best Local Similarity 96.9
Matches 312; Conservative
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96.9%; Pred. No. 7e-139;
tive 6; Mismatches 4; Indels
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Search completed: February 3, 2006, 20:46:57
Job time: 177 secs

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/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	0	UT	4.	u	2	۲.	Result No.
195	196	196	200.5	200.5	200.5	200.5	201	201	205	205	215.5	243	257.5	257.5	350	366.5	367.5	375.5	403.5	410.5	410.5	479.5	485.5	485.5	Score
11.5	11.6	11.6	11.9	11.9	11.9	11.9	11.9	11.9	12.1	12.1	12.7	14.4	15.2		20.7			22.2		24.3		28.4	28.7	28.7	Query Match 1
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US-11-127-877-57	US-11-169-976-9	US-10-502-145-1	US-10-877-346-83	US-10-055-877-340	US-10-055-877-327	US-10-055-877-248	US-10-995-561-837	US-10-995-561-838	US-10-055-877-237	US-10-055-877-225	US-11-017-058-9	US-10-980-388-97	US-10-980-388-98	US-10-980-388-39	US-10-055-877-69	US-10-055-877-247	US-10-055-877-246	US-10-980-388-119	US-10-055-877-71	US-10-055-877-245	US-10-055-877-244	US-10-980-388-113	US-10-980-388-62	US-10-980-388-111	ID
Sequence 57, Appl	Sequence 9, Appli	e 1,	83	e 34	327,	248,	837,		237,	225	Sequence 9, Appli	97,	98,		69,	247,	•		71,	245,	244,	113,	~	e 111, A	Description

Query Match Best Local Similarity

28.7%; 38.5%;

Score 485.5; DB 6; Pred. No. 3.5e-35;

Length 311;

45	44	43	42	41	40	39	38	37	36	35	34	ω ω	32	31	30	29	28	27	1
160.5	163.5	164.5	164.5	164.5	164.5	164.5	164.5	164.5	168	168.5	171	171	173	176	176	182.5	187	189.5	
9.5	9.7	9.7	9.7	9.7	9.7	9.7	9.7	9.7	9.9	10.0	10.1	10.1	10.2	10.4	10.4	10.8	11.1	11.2	1
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US-11-127-877-62	US-11-157-930-5	US-10-995-561-715	US-10-995-561-714	US-10-995-561-713	US-11-127-877-65	US-10-876-787-2	US-10-995-561-716	US-10-995-561-712	US-11-122-849-2	US-10-055-877-161	US-11-127-877-46	US-11-127-877-53	US-11-127-877-55	US-11-017-058-2	US-10-980-388-117	US-11-157-930-2	US-11-151-482-5	US-11-151-482-3	
Sequence 62, Appl	Sequence 5, Appli	Sequence 715, App	Sequence 714, App	Sequence 713, App	Sequence 65, Appl	Sequence 2, Appli	Sequence 716, App	Sequence 712, App	Sequence 2, Appli	Sequence 161, App	Sequence 46, Appl	Sequence 53, Appl	Sequence 55, Appl	Sequence 2, Appli	Sequence 117, App	Sequence 2, Appli	Sequence 5, Appli	Sequence 3, Appli	*** Add

ALIGNMENTS

US-10-980-388-111

Sequence 111, Application US/10980388 Publication No. US20050255490A1 GENERAL INFORMATION:

APPLICANT: Hiebsch, Ronald R.

APPLICANT: Hiebsch, Ronald R.

APPLICANT: Kaytes, Paul S.

APPLICANT: Word, Irinda S.

APPLICANT: Hief, Rita M.

APPLICANT: Hief, Rita M.

APPLICANT: Hief, Rita M.

APPLICANT: Hief, Rita M.

APPLICANT: Hief, Rita M.

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APPLICANT: Hief, Rita M.

APPLICANT: Hief, Rita M.

APPLICANT: Wood, Linda S.

ITILE DET INVENTION: NOWALER: US/10/980,388

CURRENT PILING DATE: 2004-11-02

PRIOR APPLICATION NUMBER: 60/184,305

PRIOR PILING DATE: 2001-02-23

PRIOR PILING DATE: 2001-02-23

PRIOR PILING DATE: 2001-02-23

PRIOR PILING DATE: 2001-02-23

PRIOR APPLICATION NUMBER: 60/184,303

PRIOR PILING DATE: 2000-02-23

PRIOR APPLICATION NUMBER: 60/184,397

PRIOR APPLICATION NUMBER: 60/184,247

PRIOR APPLICATION NUMBER: 60/184,247

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PRIOR APPLICATION NUMBER: 60/184,397

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                                                                    ; ORGANISM: Homo US-10-980-388-62
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                                                                                                    Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.0
SEQ ID NO 62
LENGTH: 530
TYPE: PRT
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Matches
              Query Match
Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR FILING DATE: 2001-02-23
PRIOR FILING DATE: 2001-02-23
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PRIOR FILING DATE: 2000-07-20
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APPLICATION NUMBER: 60/184,247
PILING DATE: 2000-02-23
CO/188,880
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APPLICATION NUMBER: 60/184,304
FILING DATE: 2000-02-23
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                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/217,369 ETILING DATE: 2000-07-11 APPLICATION NUMBER: 60/217,370 FILING DATE: 2000-07-11
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                                                                                                                                                                                                                                                                                                                           FILING DATE: 2000-03-13
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114;
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Huff, Rita M.
Wood, Linda S.
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Hiebsch, Ronald R.
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Conservative
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28.7%; Score 485.5; DB 6; 38.5%; Pred. No. 5.8e-35; tive 51; Mismatches 120;
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                                 Length 530;
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                                                                            ; ORGANISM: Homo US-10-980-388-113
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                  Query Match
Best Local Similarity
                                                                                                                                                      SOFTWARE: PatentIn version SEQ ID NO 113
    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Wood, Linda S.
TITLE OF INVENTION: Novel G Protein-Coupled
FILE REFERENCE: 00325, US1
CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
CURRENT FILING DATE: "10/04/161 633
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                                                                                                                                                                                          Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 184
                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/218,492 PRIOR FILING DATE: 2000-07-20
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                                                                                                                                LENGTH:
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Huff, Rita M.
Wood, Linda S.
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    Conservative
                  28.4%; Score 479.5; DB 6
38.2%; Pred. No. 1.1e-34;
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51;
    Mismatches 121;
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                                          DB 6;
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      Indels
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RESULT 4
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CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
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                                                       OR APPLICATION NUMBER: 60/264,117
OR FILING DATE: 2001-01-25
OR APPLICATION NUMBER: 60/264,139
OR FILING DATE: 2001-01-25
OR APPLICATION NUMBER: 60/264,478
OR FILING DATE: 2001-01-26
OR FILING DATE: 2001-01-26
             APPLICATION NUMBER: 60/263,351 FILING DATE: 2001-01-30 APPLICATION NUMBER: 60/272,870
                                                                                                                                                      APPLICATION NUMBER: 60/263,799
FILING DATE: 2001-01-24
FILING DATE:
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Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
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Patturajan, Meera
Burgess, Cahterine
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Miller, Charles
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Kekuda, Ramesh
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Spytek, Kimberly
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ısen, Bryan
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RESULT 5
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SEQ ID NO 244
LENGTH: 343
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Best Local Similarity
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PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36 IISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFSPL------PLI 88
                                                                                        SSAKPIVYFLAGRDKSQRLWEPLRVVFQRALRDGAELGEAGGSTPNTVTMEM 335
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                                                                                                                               SSANPITYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQE-TLEL 315
                                                                                                                                                                              RRORSAKLNHVILAMVSVFLVSSIYLGIDWFLF-----WVFQIPAPFPEYVTDLCICIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24.3%; Score 410.5; DB 6; 34.6%; Pred. No. 1.2e-28; tive 53; Mismatches 117;
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GENERAL INFORMATION: Sequence 245, A Publication No. APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT APPLICANT: APPLICANT APPLICANT APPLICANT: APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT: Invention: Novel Polypeptides and Nucleic Acids Encoded Thereby Padigaru, Musseller, Charles
Miller, Charles
Velizar DeCristofaro, Marc Vernet, Corine Taupier Jr., Raymond Pena, Carol Baumgartner, Jason Shimkets, Richard Gusev, Vladimir Shenoy, Suresh Burgess, Andrew, David Mezes, Peter Ratelli, Luca Kekuda, Ramesh Gerlach, Anderson, David Casman, Wolenc, Adam Eisen, Andrew Patturajan, Meera Burgess, Cahterine Zerhusen, Bryan Guo, Spytek, Kimberly Ballinger, Robert Application US/10055877 o. US20050288241A1 Xiaojia Mei Stacie David Valerie Muralidhara

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; SOFTWARE: PatentIn Ver. 2
; SEQ ID NO 245
; LENGTH: 343
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-245
                                                                                                                          Publication No. -...;
Publication No. -...;
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Miller, Charles
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PRIOR APPLICATION NUMBER: 60/272,870
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
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Gerlach, Valerie
Spytek, Kimberly
Ratelli, Luca
Kekuda, Ramesh
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                                                              Anderson, David
Ballinger, Robert
Gerlach, Valerie
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TITLE OF INVENTION: Novel Polypeptides and
FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
CURRENT FILING DATE: 2002-01-23
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PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
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PRIOR FILING DATE: 2001-01-19
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APPLICATION NUMBER: 60/272,870
FILING DATE: 2001-03-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
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                                                                                                                                                                                                    53 LLCLCGLVGNGLVLWFFGFSIKRNPFSIYFLHLASADVGYLFSKAVFSILNTGGFLGTFA 112
                                                                                                                                                                                                                                                                              36 IISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLELSGHIIFSPL-----PLI
                                                                                                                                                                                                                                                                                                                            Similarity
SANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQE-TLEL
                                                                                KMPLT-RLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDW--KVLFCHVHLVSIFLSALN
                                                                                                                                                                                  DYIRSVCRVLGLCM---FLTGVSLLPAVSAERCASVIFPAWYWRRPKRLSAVVCALLWV
                                                  RRQRSAKLNHVÍLAMVSVÉLVSSIYLGÍDWELE-----WVFQIPAPEPEYVTDLCICIN
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Shimkets, Richard
Gusev, Vladimir
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Taupier Jr., Raymond
Pena, Carol
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Patturajan, Meera
Burgess, Cahterine
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Eisen, Andrew
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Andrew, David
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usen, Bryan
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                                                                                                                                                                                                                                                                                                             23.9%; Score 403.5; DB 6
34.2%; Pred. No. 4.7e-28;
ative 53; Mismatches 118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.0
SEQ ID NO 119
LENGTH: 323
TYPE: PRT
ORGANISM: Homo sapiens
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TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference To Related.

FILE REFERENCE: 00325.US1

CURRENT APPLICATION NUMBER: US/10/980,388

CURRENT FILING DATE: 2004-01-02

PRIOR APPLICATION NUMBER: US/09/791,932

PRIOR FILING DATE: 2001-02-23

PRIOR APPLICATION NUMBER: 60/184,305

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR FILING DATE: 2000-02-23

PRIOR FILING DATE: 2000-02-23

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PRIOR FILING DATE: 2000-02-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 102;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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PRIOR APPLICATION NUMBER: 60/217,370
PRIOR FILING DATE: 2000-07-11
PRIOR APPLICATION NUMBER: 60/218,492
PRIOR FILING DATE: 2000-07-20
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NUMBER OF SEQ ID NOS: 184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR
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APPLICATION NUMBER: 60/188,880
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APPLICATION NUMBER: 60/184,247
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    278
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                                                                                                                                                                                                                                 152 LRSILEWMFCDFLFSGANSVWCETSDFITIAWLVFLCVVLCGSSLVLLVRILCGSRKMPL 211
                                                                                                                                                                                                                                                                                                                                                                                                                   33 LTCIISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFSPLPLINIRH 92
                                                                                                                                                                                                                                                                                                                                                                              52
                                                                                                                                                                                                                                                                                                                           93 PISKIĻSPYMTFPYF-IGLSMĻSAISTERCLSILWPIWYHCRRPRYLSSVMCVLLWALSL 151
                                                                                                                                                                                                                                                                               GAQDTLYEVLTELMEAVGLMLLAAFSVERCLSDLFPACYQGCRERHASAVLCALVMTPTL 168
SSKPLIYSGLG--ROPGKREPLRSVLRRALGEGAELGARGOSLPMGLL 323
                                         SANPITYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQETL 313
                                                                                            PRLYGIVLGALLLLFFCGLPSVFYWSLQPLLNF--
                                                                                                                                      TRLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDWKVLFCHVHLVSIF-----LSALNS 265
                                                                                                                                                                                       PAVPLPANACGLLRNSACPLVCPRYHVASVTWFLVLARVAWTAGVVLFVWVTCCSTR-PR
                                                                                                                                                                                                                                                                                                                                                                          LTLIVGLGGPVGNGLVLWNLGFRIKKGPFSIYLLHLAAADFLFLSCRVGFS---VAQAAL 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ruff, Valerie
Huff, Rita M.
Wood, Linda S.
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                                                                                            -LLPVFSPLATLLACVNS
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) ORGANISM: Homo sapiens
US-10-055-877-246
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US-10-055-877-246
Query Match
Best Local Similarity
Matches 96; Conserv
                                                                                                              SOFTWARE: Patentin Ver.
SEQ ID NO 246
LENGTH: 342
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                                                                                                                                                                  Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 512
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PRIOR APPLICATION NUMBER: 60/263,598
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CURRENT FILING DATE: 2002-01-22
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PRIOR APPLICATION NUMBER: 60/264,117
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PRIOR APPLICATION NUMBER: 60/263,799
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Miller, Charles
Miller, Velizar
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Mezes, Peter
Patturajan, Meera
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Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
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Wolenc, Adam
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Kekuda, Ramesh
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Spytek, Kimberly
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Gerlach, Valerie
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   Conservative
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 21.7%; Score 367.5; DB 6; 32.4%; Pred. No. 6.1e-25; ative 61; Mismatches 110;
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   Indels
                                Length 342;
   29;
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                                                                                                                     PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
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TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
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           OR APPLICATION NUMBER: 60/264,117
OR FILING DATE: 2001-01-25
OR APPLICATION NUMBER: 60/264,139
OR FILING DATE: 2001-01-25
OR APPLICATION NUMBER: 60/264,478
OR APPLICATION NUMBER: 60/264,478
OR APPLICATION NUMBER: 60/263,351
OR APPLICATION NUMBER: 60/263,351
OR FILING DATE: 2001-01-30
APPLICATION
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Miller, Charles
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Shimkets, Richard
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Mezes, Peter
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Kekuda, Ramesh
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Spytek, Kimberly
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o. US20050288241A1
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NUMBER: 60/272,870
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; LENGTH: 319
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PRIOR APPLICATION NUMBER: 60/275,990
PRIOR PILING DATE: 2001-03-14
PRIOR PILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 512
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 247
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Best Local Similarity
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261 SALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQB-TLEL 315
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                                                                                                                                                                                                                                                                                                                                 36 IISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFSPLPLIN-----
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                                                                                                         CGSRKMPLT-RLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDW--KVLFCHVHLVSIFL
                                                                                                                                                                                 LIWALSLIRSILEWMFCDFLFSGANSVWCETSDFITIAWLVFL-CVVLCGSSLVLLVRIL 203
                                                                                                                                                                                                                                                        ----IRHPISKILSPVMTFPYFIGLSMLSAISTERCLSILWPIWYHCRRPRYLSSVMCV 144
                                                                                                                                               LLWMLSFLVTSIHNYFCMFLGHEAPGTVCRNMDIALGILLFFLFCPLMVLPCLALILHVE
                                                                      CRARRORSAKLNHVVLAIVSVFLVSSIYLGIDWFLF-----WVFQIPAPFPEYVTDLC
                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 musculus
                                                                                                                                                                                                                                                                                                                                                                                      21.7%; Score 366.5; DB 6 33.1%; Pred. No. 6.9e-25;
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US-10-055-877-69 RESULT 10 GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc Sequence 69, Application US/10055877 Publication No. US20050288241A1 APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT APPLICANT: APPLICANT APPLICANT: APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT: APPLICANT APPLICANT APPLICANT APPLICANT: Gerlach, Vares, Spytek, Kimberly Ratelli, Luca Keuda, Ramesh Guo, Xiaojia Li, Li Casman, Padigaru, Muraii Miller, Charles Vernet, Corine Taupier Jr., Raymond Pena, Carol Boldog, Shenoy, Suresh Gusev, Vladimir Baumgartner, Jason Shimkets, Richard Wolenc, Adam Eisen, Andrew Mezes, Peter Patturajan, Meera Burgess, Cahterine Zerhusen, Bry Andrew, David Ballinger, Anderson, David Zhong, Mei Ference Stacie Muralidhara Robert

OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby

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US-10-980-388-39
; Sequence 39, Application US/10980388
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Best Local Similarity
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APPLICANT: Parod
APPLICANT: Hiebs
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NUMBER OF SEQ ID NOS: 512
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PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
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CURRENT FILING DATE: 2002-01-22
                   TITLE OF
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REFERENCE: 00325.US1
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APPLICATION NUMBER: 60/264,139
FILING DATE: 2001-01-25
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                                                                                                                                                                                         INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , 68
                 INVENTION: Novel G Protein-Coupled Receptors Cross-Reference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LLCLCGLVGNGLVLWFFGFSIKRNPFSIYFLHLASADVGYLFSKAVFSILNTGGFLGTFA 112
                                                                                                                                                                                                                                                                                                                                                            SANPITYFFVGSFRORQNRQNLKLVLQRA 294
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LSLLRSILEWMFCDFLFSGANSVWCETSD-FITIAWLVFLCVVLCGSSLVLLVRILCGSR 207
                                                                                                                                                                                                                                                                                                                         SAKPIVYFLAGR-TSRSGCWSLRVVFSGA
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                                                                                                            Parodi, Luis A.
Hiebsch, Ronald R.
Lind, Peter
                                                               Ruff, Valerie
                                                                                        Kaytes,
                                 Huff, Rita M. Wood, Linda S.
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                                                                                                                                                                     , Gabriel
                                                                                          Paul S.
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Pred. No. 2e-23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20;
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                 To
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               Related App
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APPLICANT: Vogeli,
APPLICANT: Parodi,
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APPLICANT:
APPLICANT:
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Best Local
            PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 60/184,305
PRIOR FILING DATE: 2000-02-23
PRIOR APPLICATION NUMBER: 60/184,304
PRIOR FILING DATE: 2000-02-23
PRIOR APPLICATION NUMBER: 60/184,303
                                                                                                                                                     TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference To Related Appli
FILE REFERENCE: 00325.US1
CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
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TYPE: PRT
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CURRENT FILING DATE: 2004-11-02
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FILING DATE: 2000-02-23
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Lind, Peter
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RUEE, Paul S.

RUEE, Valerie

LANT: Huff, Valerie

TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference To Related Appl

CURRENT APPLICATION NUMBER: US/10/980,388

CURRENT APPLICATION NUMBER: US/10/980,388

CURRENT FILING DATE: 2004-11-02

PRIOR APPLICATION NUMBER: US/9/791,932

PRIOR APPLICATION NUMBER: 60/184,305

PRIOR RILING DATE: 2001-02-23

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR APPLICATION NUMBER: 60/184,304

PRIOR APPLICATION NUMBER: 60/187

PRIOR FILING DATE: 2000-02-23

PRIOR APPLICATION NUMBER: 60/187

PRIOR FILING DATE: 2000-02-23

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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-980-388-98
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SEQ ID NO 98
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PRIOR APPLICATION NUMBER: 60/188,880
PRIOR FILING DATE: 2000-03-13
PRIOR APPLICATION NUMBER: 60/217,369
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OR FILING DATE: 2000-02-23
OR APPLICATION NUMBER: 60/184,397
OR FILING DATE: 2000-02-23
OR APPLICATION NUMBER: 60/184,247
OR APPLICATION NUMBER: 60/184,880
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OR APPLICATION NUMBER: 60/188,880
OR FILING DATE: 2000-03-13
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les 57; Conserv
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APPLICATION NUMBER: 60/184,247
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APPLICATION NUMBER: 60/217,370
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                           APPLICATION NUMBER: 60/217,369
FILING DATE: 2000-07-11
APPLICATION NUMBER: 60/217,370
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FILING DATE: 2000-07-11
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; LENGTH: 211
; TYPE: PRT
; ORGANISM: Homo 8
US-10-980-388-97
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CURRENT FILING DATE: 2004-12-21
PRIOR APPLICATION NUMBER: US 09/101,518
PRIOR FILING DATE: 1998-12-21
PRIOR APPLICATION NUMBER: PCT/US96/00499
PRIOR FILING DATE: 1996-01-11
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
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NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.0
SEQ ID NO 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9, Application US/11017058
Publication No. US20060014243A1
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Best Local Similarity
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PRIOR FILING DATE: 2000-07-20
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TITLE OF INVENTION: Human G-Protein Chemokine
FILE REFERENCE: PF218C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 353
TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              109 GAQDTLYFVLTFLWFAVGLWLLAAFSVERCLSDLFPACYQGCRPRHASAVLCALVWTPTL
                                                                         195 GNNTANWRMLLRILPQSFGFIVPLLIMLFCYGFTLRTLFKAHMGQKHRAMRVIFAVVLIF
                                                                                                                                                        140
                                                                                                                                                                                         122 LSILWPIWYHCRR----PRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWC-ETS 176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93
                                                                                                                                                                                                                                  89 LLFALTLPIWAASKVNGWIFGTFL-----CKVVSLLKEVNFYSGILLLACISVDRY 139
                                                                                                                                                                                                                                                                      73 FLF-----LSGHIIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTERC 121
                                                                                                                                                                                                                                                                                                             29 DAAPĆEPESLBINKYFVVIIYALVFLLSLIGNSLVMLVILYSRVGRSVTDVYLLNLALAD 88
                                                                                                                                                                                                                                                                                                                                            19 EETPCYNQTLSFTG-----LTCIISLVALTGNA-VVLWLLGCRMRRNAVSIYILNLVAAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ch 14.4%; Score 243; DB 6; 1 Similarity 38.4%; Pred. No. 2.2e-14; 61; Conservative 26; Mismatches 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     353
                                                                                                                                                                                                                                                                                                                                                                                             81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PISKILSPYMTEPYF-IGLSMLSAISTERCLSILWPIWYHCRRPRYLSSYMCVLLWALSL 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PAVPLPANACGLIRNSACPLVCPRYHVASVTWFLVLARV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LRSILEWMFCDFLFSGANSVWCETSDFITIAWLVFLCVV 190
                                                                                                                DFITIAMLVFLCVV--LCGSSLVLLVRILC-GSRKMPLTRLYV------TILLTVLVF
                                                                                                                                                      LAIV-----HATRTLTQKRYLVKFICLSIWGLSLLLALPVLLFRRTVYSSNVSPACYEDM 194
LLCWLPYNLVLLADTLMRTQVIQETCERRNHIDRALDATEI-----LGILHSCLNPLIY
                                      LLCGLPFG-----
                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                           12.7%; Score 215.5; DB 7; 26.3%; Pred. No. 8.7e-12; ative 55; Mismatches 105;
                                    IQWALFSRIHLDWKVLFCHVHLVSIFLSALNSSANPIIY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             68; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Receptor HSATU68
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 353;
                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                   67;
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        308
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RESULT 15 US-10-055-877-225 유 성

273 FFVG-SFR 279 |:| || 309 AFIGQKFR 316

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PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-03-02
PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
                                    Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 512
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 225
LENGTH: 259
TYPE: PAT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR FILING DATE: 2001-01-19
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APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
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OTHER INFORMATION: Description of Artificial Sequence: 7tm_1 domain
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                         FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shenoy, Suresh
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gusev, Vladimir
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Shimkets, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Burgess, Cahterine
Eisen, Andrew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mezes, Peter
Patturajan, Meera
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Zerhusen, Bryan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ratelli, Luca
Kekuda, Ramesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gerlach, Valerie
Spytek, Kimberly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anderson, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zhong, Mei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wolenc, Adam
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b. US20050288241A1
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                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: consensus sequence US-10-055-877-225
                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
226 IMSSTCELERVLPTALLVTLWLAYVNSCLNPIIY
                                                                    172 GFLLPLLVILVCYTRILRTLRKAAKTLLVVVVVVFVLCWLPY-----FIVLLLDTLCLSI 225
                                                                                                     193 GSSLVILLVRILCGSRKM-PLTRLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDWKVL--
                                                                                                                                         117 LLSL-----PPLLFSWVKTVEEGNGTLNVNVTVCLIDFPEESTASVSTWLRSYVLLSTLV 171
                                                                                                                                                                          152 LRSILEWMFCDFLFSGANSV---------WCETSDFITIAWL--VFLCVVLC 192
                                                                                                                                                                                                          58 LCKLVTALDVVNMYASILLITAISIDRYLAIVHPLRYRRRRTSPR-RAKVVILLVMVLAL 116
                                                                                                                                                                                                                                               94 ISKILSPVMTFPYFIGLSMLSAISTERCLSILWPIWYHCRR--PRYLSSVMCVLLWALSL
                                                                                                                                                                                                                                                                                                                   44 GN-AVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFSPLPLINI-----RHP
                                                                                                                                                                                                                                                                               80; Conservative
                                ----FCHVH-----LVSIFLSALNSSANPITY 272
                                                                                                                                                                                                                                                                                                                                                                      12.1%;
                                                                                                                                                                                                                                                                                                                                                    Score 205; DB 6;
Pred. No. 5.2e-11;
6; Mismatches 88
 259
                                                                                                                                                                                                                                                                                                                                                                                     Length 259;
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                                                                                                                                                                                                                                                                                                                                                      60;
                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                       249
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Search completed: February 3, 2006, 20:47:18
Job time : 17 secs

Ipis bade Blank (nabto)

November 2005

Published_Applications Nucleic Acid and Published_Applications Amino Acid database searches now generate two sets of results each. The Published_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published_Applications_New databases; older published applications make up the Published_Applications_Main databases.

Searches run against Nucleic Acid Published_Applications produce two sets of results, with the extensions '.rnpbm (Published_Applications_NA_New).

Searches run against Amino Acid Published_Applications produce two sets of results, with the extensions .rapbm (Published_Applications_AA_New).

IVI? Lage Blank (uspto)